

(12) INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

(19) World Intellectual Property
Organization
International Bureau



(43) International Publication Date
4 November 2004 (04.11.2004)

PCT

(10) International Publication Number
WO 2004/093902 A1

(51) International Patent Classification⁷: **A61K 38/29**,
C07K 1/06, 14/635

Boston, MA 02114 (US). **SHIMIZU, Naoto** [JP/US];
Pharmaceutical Research Dept. II, Chugai Pharmaceutical
Co., Ltd, 1-135 Komakado, Gotemba, Shizuoka 412-8513
(JP). **CARTER, Percy** [US/US]; 34 Parkside Drive,
Princeton, NJ 08504 (US).

(21) International Application Number:
PCT/US2003/008261

(22) International Filing Date: 19 March 2003 (19.03.2003)

(74) Agents: **BUGAISKY, Lawrence, B.** et al.; Sterne,
Kessler, Goldstein & Fox P.L.L.C., 1100 New York Av-
enue, N.W., Washington, DC 20005-3934 (US).

(25) Filing Language: English

(26) Publication Language: English

(81) Designated States (*national*): AE, AG, AL, AM, AT, AU,
AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CO, CR, CU,
CZ, DE, DK, DM, DZ, EC, EE, ES, FI, GB, GD, GE, GH,
GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC,
LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW,
MX, MZ, NO, NZ, OM, PI, PL, PT, RO, RU, SC, SD, SE,
SG, SK, SL, TJ, TM, TN, TR, TT, TZ, UA, UG, US, UZ,
VC, VN, YU, ZA, ZM, ZW.

(71) Applicants (*for all designated States except US*): **THE
GENERAL HOSPITAL CORPORATION** [US/US];
Fruit Street, Boston, MA 02492 (US). **BRISTOL-MY-
ERS SQUIBB COMPANY** [US/US]; Route 206 &
Provinceline Road, Princeton, NJ 08543-4000 (US).

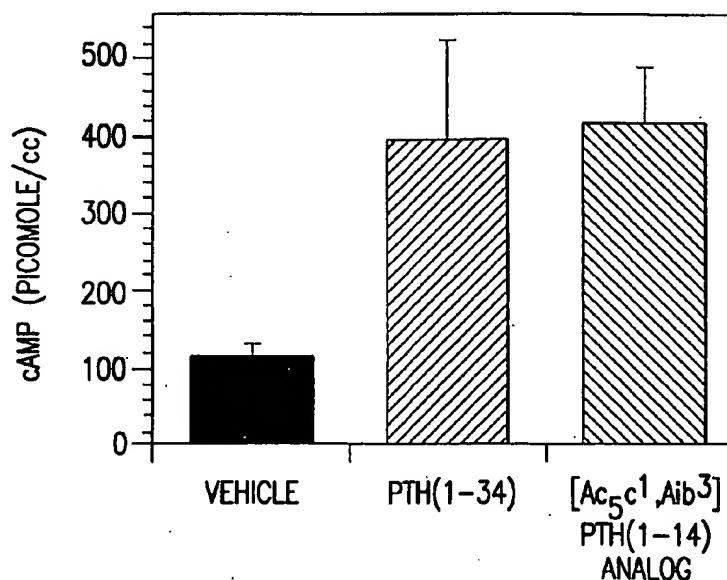
(72) Inventors; and

(75) Inventors/Applicants (*for US only*): **GARDELLA,
Thomas, J.** [US/US]; 1045 Greendale Avenue, Needham,
MA 02492 (US). **POTTS, John, T., Jr.** [US/US]; Building
149, 13th Street, Charlestown, MA 02129 (US). **KRO-
NENBERG, Henry, M.** [US/US]; 50 Blossom Street,

(84) Designated States (*regional*): ARIPO patent (GH, GM,
KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW),
Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM),
European patent (AT, BE, BG, CH, CY, CZ, DE, DK, EE,
ES, FI, FR, GB, GR, HU, IE, IT, LU, MC, NL, PT, RO,

[Continued on next page]

(54) Title: CONFORMATIONALLY CONSTRAINED PARATHYROID HORMONES WITH α -HELIX STABILIZERS



(57) Abstract: The present invention relates to conformationally constrained parathyroid hormone (PTH) analogs and derivatives of those analogs. The invention also provides methods of preparing and using the PTH analogs. Further, the invention encompasses compositions and methods for use in limiting undesired bone loss in a vertebrate at risk of such bone loss, in treating conditions that are characterized by undesired bone loss or by the need for bone growth, e.g. in treating fractures or cartilage disorders and for raising camp levels in cells where deemed necessary.

Best Available Copy

WO 2004/093902 A1



SE, SI, SK, TR), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG).

— with a declaration as to non-prejudicial disclosures or exceptions to lack of novelty

Published:

— with international search report

For two-letter codes and other abbreviations, refer to the "Guidance Notes on Codes and Abbreviations" appearing at the beginning of each regular issue of the PCT Gazette.

Conformationally Constrained Parathyroid Hormones With α -Helix Stabilizers

5 STATEMENT REGARDING FEDERALLY-SPONSORED RESEARCH AND DEVELOPMENT

Statement under MPEP 310. The U.S. Government has a paid-up license
in this invention and the right in limited circumstances to require the patent
10 owner to license others on reasonable terms as provided for by the terms of Grant
No. DK-11794 awarded by the National Institute of Health.

Background of the Invention

Field of the Invention

15 The present invention relates to conformationally constrained parathyroid
hormone (PTH) analogs, and methods of preparing and using the PTH analogs.

Background Art

Parathyroid hormone

Parathyroid hormone (PTH), an 84 amino acid peptide, is the principal
20 regulator of ionized blood calcium in the human body (Kronenberg, H.M., *et al.*,
In *Handbook of Experimental Pharmacology*, Mundy, G.R., and Martin, T.J.,
(eds), pp. 185-201, Springer-Verlag, Heidelberg (1993)). Regulation of calcium
concentration is necessary for the normal function of the gastrointestinal, skeletal,
neurologic, neuromuscular, and cardiovascular systems. PTH synthesis and
25 release are controlled principally by the serum calcium level; a low level
stimulates and a high level suppresses both hormone synthesis and release. PTH,
in turn, maintains the serum calcium level by directly or indirectly promoting
calcium entry into the blood at three sites of calcium exchange: gut, bone, and
kidney. PTH contributes to net gastrointestinal absorption of calcium by favoring

-2-

the renal synthesis of the active form of vitamin D. PTH promotes calcium resorption from bone indirectly by stimulating differentiation of the bone-resorbing cells, osteoclasts. It also mediates at least three main effects on the kidney: stimulation of tubular calcium reabsorption, enhancement of phosphate clearance, and promotion of an increase in the enzyme that completes synthesis of the active form of vitamin D. PTH is thought to exert these effects primarily through receptor-mediated activation of adenylate cyclase and/or phospholipase C.

Disruption of calcium homeostasis may produce many clinical disorders (*e.g.*, severe bone disease, anemia, renal impairment, ulcers, myopathy, and neuropathy) and usually results from conditions that produce an alteration in the level of parathyroid hormone. Hypercalcemia is a condition that is characterized by an elevation in the serum calcium level. It is often associated with primary hyperparathyroidism in which an excess of PTH production occurs as a result of a parathyroid gland lesion (*e.g.*, adenoma, hyperplasia, or carcinoma). Another type of hypercalcemia, humoral hypercalcemia of malignancy (HHM) is the most common paraneoplastic syndrome. It appears to result in most instances from the production by tumors (*e.g.*, squamous, renal, ovarian, or bladder carcinomas) of a class of protein hormone which shares amino acid homology with PTH. These PTH-related proteins (PTHrP) appear to mimic certain of the renal and skeletal actions of PTH and are believed to interact with the PTH receptor in these tissues.

Osteoporosis

Osteoporosis is a potentially crippling skeletal disease observed in a substantial portion of the senior adult population, in pregnant women and even in juveniles. The term osteoporosis refers to a heterogeneous group of disorders. Clinically, osteoporosis is separated into type I and type II. Type I osteoporosis occurs predominantly in middle aged women and is associated with estrogen loss at menopause, while osteoporosis type II is associated with advancing age. Patients with osteoporosis would benefit from new therapies designed to promote

fracture repair, or from therapies designed to prevent or lessen the fractures associated with the disease.

The disease is marked by diminished bone mass, decreased bone mineral density (BMD), decreased bone strength and an increased risk of bone fracture. At present, there is no effective cure for osteoporosis, though estrogen, calcitonin and the bisphosphonates, etidronate and alendronate are used to treat the disease with varying levels of success. These agents act to decrease bone resorption. Since parathyroid hormone regulates blood calcium and the phosphate levels, and has potent anabolic (bone-forming) effects on the skeleton, in animals (Shen, V., *et al.*, *Calcif. Tissue Int.* 50:214-220 (1992); Whitefield, J.F., *et al.*, *Calcif. Tissue Int.* 56:227-231 (1995) and Whitfield, J.F., *et al.*, *Calcif. Tissue Int.* 60:26-29 (1997)) and humans (Slovik, D.M., *et al.*, *J. Bone Miner. Res.* 1:377-381 (1986); Dempster, D.W., *et al.*, *Endocr. Rev.* 14:690-709 (1993) and Dempster, D.W., *et al.*, *Endocr. Rev.* 15:261 (1994)) when administered intermittently, PTH, or PTH derivatives, are prime candidates for new and effective therapies for osteoporosis.

PTH Derivatives

PTH derivatives include polypeptides that have amino acid substitutions or are truncated relative to the full length molecule. Both a 14 and a 34 amino acid amino-terminal truncated form of PTH, as well as a C-terminal truncated form have been studied. Additionally, amino acid substitutions within the truncated polypeptides have also been investigated.

Synthetic PTH(1-34) exhibits full bioactivity in most cell-based assay systems, has potent anabolic effects on bone mass in animals and has been shown to reduce the risk of bone fracture in postmenopausal osteoporotic women (Neer, R.M., *et al.*, *N.E.J.M.* 344:1434-1441 (2001); Dempster, D.W., *et al.*, *Endocr Rev* 14:690-709 (1993)). PTH acts on the PTH/PTHrP receptor (P1R), a class II G protein-coupled heptahelical receptor that couples to the adenylyl cyclase/cAMP and phospholipase C/inositol phosphate (IP) signaling pathway (Rippner, H., *et al.*, *Science* 254:1024-1026 (1991)). Deletion analysis studies have shown that the amino-terminal residues of PTH play a crucial role in stimulating the P1R to

-4-

activate the cAMP and IP signaling pathways (Tregear, G.W., *et al.*, *Endocrinology* 93:1349-1353 (1973); Takasu, H., *et al.*, *Biochemistry* 38:13453-13460 (1999)). Crosslinking and receptor mutagenesis studies have indicated that residues in the amino-terminal portion of PTH interact with the extracellular loops and extracellular ends of the seven transmembrane helices, which reside within the juxtamembrane region of the receptor (Bergwitz, C., *et al.*, *J. Biol. Chem.* 271:26469-26472 (1996); Hoare, S.R.J., *et al.*, *J. Biol. Chem.* 276:7741-7753 (2001); Behar, V., *et al.*, *J. Biol. Chem.* 275:9-17 (1999); Shimizu, M., *et al.*, *J. Biol. Chem.* 275:19456-19460 (2000); Luck, M.D., *et al.*, *Molecular Endocrinology* 13:670-680 (1999)).

α -Helix Stabilizers

The first 34 amino acids of PTH and PTHrP contain sufficient information for high affinity P1R binding and potent induction of P1R-mediated signaling responses (Neer, RM, *et al.*, *N.E.J.M.* 344: 1434-1441 (2001)). Short N-terminal fragments of PTH, such as PTH(1-14) and PTH(1-11) exhibit extremely weak binding affinities ($K_d \gg 100 \mu\text{M}$) but are nonetheless capable of eliciting cAMP-signaling responses, albeit with potencies ($\text{EC}_{50} \geq 100 \mu\text{M}$) that are substantially weaker than that of PTH(1-34) ($\text{EC}_{50} \sim 2 \text{ nM}$) (Luck, MD *et al.*, *Molecular Endocrinology* 13: 670-680 (1999)). It has been reported that a series of modified PTH(1-14) and PTH(1-11) analogs exhibit signaling potencies that are nearly, or even fully, equal to that of PTH(1-34) (Shimizu, M *et al.*, *Endocrinology* 142: 3068-3074 (2001); Shimizu, M. *et al.*, *J. Biol. Chem.* 276: 490003-49012 (2001); Shimizu, M. *et al.*, *J. Biol. Chem.* 275: 21836-21843 (2000)).

Recently, it was also reported that PTH(1-14) analogs containing the α , α -disubstituted amino acid, α -aminoisobutyric acid (Aib) at positions 1 and/or 3, have 10- to 100- fold higher affinities and cAMP signaling potencies than do their counterpart peptides containing alanine at these positions (Shimizu, N. *et al.* *J. Biol. Chem.* 276: 49003-49012 (2001)).

Brief Summary of the Invention

The invention provides novel PTH polypeptide derivatives containing amino acid substitutions at selected positions in the polypeptides. The derivatives function as full, or nearly full, agonists of the PTH-1 receptor. Because of their unique properties, these polypeptides have utility as drugs for treating human diseases of the skeleton, such as osteoporosis.

The invention provides derivatives of PTH(1-14), PTH(1-13), PTH(1-12), PTH(1-11), PTH(1-10) and PTH(1-9) polypeptides, wherein at least one residue in each polypeptide is an α,α -di-substituted amino acid. These polypeptides may also contain a residue which is a helix, preferably an α -helix, stabilizing residue. These α -helix stabilizing residues include, but are not limited to α,α -di-alkyl amino acids with structurally varied sidechains, such as: α -aminoisobutyric acid (Aib), α,α -diethyl-glycine (Deg), 1-aminocyclopropane-1-carboxylic acid (Ac_3c), 1-aminocyclopentane-1-carboxylic acid (Ac_5c), amino-cyclobutane-1-carboxylic acid (Ac_4c), and 1-amino-cyclohexane-1-carboxylic acid (Ac_6c). The invention also provides methods of making such peptides. Further, the invention encompasses compositions and methods for use in limiting undesired bone loss in a vertebrate at risk of such bone loss, in treating conditions that are characterized by undesired bone loss or by the need for bone growth, e.g. in treating fractures or cartilage disorders and for raising cAMP levels in cells where deemed necessary.

In one aspect, the invention is directed to a biologically active peptide consisting essentially of $X_{01}ValX_{02}GluIleGlnLeuMetHisX_{03}X_{04}X_{05}X_{06}X_{07}(SEQ. ID. NO.1)$, wherein X_{01} is an α -helix-stabilizing residue, Gly, Ser or Ala; X_{02} is an α -helix-stabilizing residue, Ala or Ser; X_{03} is Ala, Gln or Asn; X_{04} is Arg, Har or Leu; X_{05} is an α -helix stabilizing residue, Ala or Gly; X_{06} is an α -helix stabilizing residue or Lys; X_{07} is an α -helix stabilizing residue, Trp or His; wherein at least one of X_{01} , X_{02} , X_{03} , X_{04} , X_{05} , X_{06} or X_{07} is an α -helix stabilizing residue, and wherein at least one of said α -helix stabilizing residues is Aib (α -aminoisobutyric acid), Ac_3c (1-aminocyclopropane-1-carboxylic acid), Ac_4c (1-

-6-

amino-cyclobutane-1-carboxylic acid), Ac₃c (1-aminocyclopentane-1-carboxylic acid), Ac₆c (1-amino-cyclohexane-1-carboxylic acid), or Deg (α,α -diethylglycine).

The invention is further drawn to fragments of the peptide having the sequence of SEQ. ID. NO. 1, in particular

5 X₀₁ValX₀₂GluIleGlnLeuMetHisX₀₃X₀₄X₀₅X₀₆ (part of SEQ. ID. NO. 1),
 X₀₁ValX₀₂GluIleGlnLeuMetHisX₀₃X₀₄X₀₅ (part of SEQ. ID. NO. 1),
 X₀₁ValX₀₂GluIleGlnLeuMetHisX₀₃X₀₄ (part of SEQ. ID. NO. 1),
 X₀₁ValX₀₂GluIleGlnLeuMetHisX₀₃ (part of SEQ. ID. NO. 1), and
 X₀₁ValX₀₂GluIleGlnLeuMetHis (part of SEQ. ID. NO. 1). The invention further

10 encompasses pharmaceutically acceptable salts of the above-described peptides, and N- or C-derivatives of the peptides, wherein at least one of X₀₁, X₀₂, X₀₃, X₀₄, X₀₅, or X₀₆ is an α -helix-stabilizing residue. One or more of the α -helix-stabilizing residue may be selected from the group consisting of Aib, Ac₃c, Ac₄c, Ac₅c, Ac₆c, or Deg.

15 In addition, the invention is drawn to a biologically active polypeptide consisting essentially of X₀₁ValX₀₂GluIleGlnLeuMetHisGlnHarAlaLysTrp (SEQ. ID. NO. 2) as well as fragment peptides containing amino acids 1-13, 1-12, 1-11, 1-10, or 1-9. The invention further encompasses pharmaceutically acceptable salts of the above-described peptides and N- or C-derivatives of the peptides,

20 wherein X₀₁ is an α -helix-stabilizing residue, Gly, Ser or Ala; and X₀₂ is an α -helix-stabilizing residue, Ala, or Ser. α -helix stabilizing residues include, but are not limited to, the group consisting of Aib, Ac₃c, Ac₄c, Ac₅c, Ac₆c, and Deg as defined above.

Preferred embodiments of the biologically active peptide include:

25 Ac₅cValAibGluIleGlnLeuMetHisGlnArgAlaLysTrpNH₂ (SEQ.ID.NO. 30),
 Ac₅cValAlaGluIleGlnLeuMetHisGlnHarAlaLysTrpNH₂ (SEQ. ID. NO. 4),
 AibValAc₅cGluIleGlnLeuMetHisAsnLeuGlyLysHisNH₂(SEQ. ID. NO. 32), and
 Ac₅cValAibGluIleGlnLeuMetHisNH₂ (SEQ. ID. NO. 31). It is contemplated that
 fragments of the above mentioned peptides, containing amino acids 1-9, 1-10, 1-
 30 11, 1-12 or 1-13, are also embodiments of the present invention. The invention

-7-

further encompasses pharmaceutically acceptable salts of the above-described peptides, and N- or C-derivatives of the peptides.

In another aspect, the invention is directed to a biologically active peptide consisting essentially of X_{01} Val X_{02} GluIle X_{03} LeuMetHis X_{04} X_{05} X_{06} Lys X_{07} (SEQ. ID. NO. 5), wherein X_{01} is an α -helix-stabilizing residue, Gly, Ser or Ala; X_{02} is an α -helix-stabilizing residue, Ala or Ser; X_{03} is Ala, Gln or Asn; X_{04} is Ala, Gln or Asn; X_{05} is an α -helix-stabilizing residue, Ala, Gly, Har or Arg; X_{06} is an α -helix-stabilizing residue or Lys; and X_{07} is an α -helix-stabilizing residue, Trp or His; wherein at least one of X_{01} , X_{02} , X_{03} , X_{04} , X_{05} , X_{06} or X_{07} is an α -helix-stabilizing residue, and wherein at least one of said α -helix stabilizing residues is Aib, Ac₃c, Ac₄c, Ac₅c, Ac₆c, or Deg.

The invention is further drawn to fragments of the peptide having the sequence of SEQ. ID. NO. 5, in particular X_{01} Val X_{02} GluIle X_{03} LeuMetHis X_{04} X_{05} X_{06} Lys (part of SEQ. ID. NO. 5), X_{01} Val X_{02} GluIle X_{03} LeuMetHis X_{04} X_{05} X_{06} (part of SEQ. ID. NO. 5), X_{01} Val X_{02} GluIle X_{03} LeuMetHis X_{04} X_{05} (part of SEQ. ID. NO. 5), X_{01} Val X_{02} GluIle X_{03} LeuMetHis X_{04} (part of SEQ. ID. NO. 5), and X_{01} Val X_{02} GluIle X_{03} LeuMetHis (part of SEQ. ID. NO. 5), wherein X_{01} is an α -helix-stabilizing residue, Gly, Ser or Ala; X_{02} is an α -helix-stabilizing residue, Ala, or Ser; X_{03} is Ala, Gln or Asn; X_{04} is Ala, Gln or Asn; X_{05} is an α -helix-stabilizing residue, Ala, Gly, Har or Arg; X_{06} is an α -helix-stabilizing residue or Lys; and X_{07} is an α -helix-stabilizing residue, Trp or His; wherein at least one of X_{01} , X_{02} , X_{03} , X_{04} , X_{05} , X_{06} or X_{07} is an α -helix-stabilizing residue, and wherein at least one of said α -helix stabilizing residues is Aib, Ac₃c, Ac₄c, Ac₅c, Ac₆c, or Deg. The invention further encompasses pharmaceutically acceptable salts of the above-described peptides, and N- or C- derivatives of the peptides.

In addition, the invention is drawn to a biologically active polypeptide consisting essentially of X_{01} Val X_{02} GluIleGlnLeuMetHisGlnHarAlaLysTrp-amide (SEQ. ID. NO. 6) and fragment peptides containing amino acids 1-13, 1-12, 1-11, 1-10, or 1-9. The invention further encompasses pharmaceutically acceptable salts of the above-described peptides, and N- or C-derivatives of the peptides, wherein

-8-

X₀₁ is an α -helix-stabilizing residue, Gly, Ser or Ala; and X₀₂ is an α -helix-stabilizing residue, Ala or Ser, and wherein at least one of said α -helix stabilizing residues is Aib, Ac₃c, Ac₄c, Ac₅c, Ac₆c, or Deg.

Preferred embodiments of the biologically active peptide include:

5 Ac₄cValAibGluIleGlnLeuMetHisGlnHarAlaLysTrp (SEQ. ID. NO. 7),
Ac₆cValAibGluIleGlnLeuMetHisGlnHarAlaLysTrp (SEQ. ID. NO. 8), Ac₅cVal
Ac₄cGluIleGlnLeuMetHisGlnHarAlaLysTrp (SEQ. ID. NO. 9),
Ac₅cValAc₆cGluIleGlnLeuMetHisGlnHarAlaLysTrp (SEQ. ID. NO. 10), Ac₄cVal
Ac₄cGluIleGlnLeuMetHisGlnHarAlaLysTrp (SEQ. ID. NO. 11), and Ac₆cVal
10 Ac₆cGluIleGlnLeuMetHisGlnHarAlaLysTrp (SEQ. ID. NO. 12). It is
contemplated that fragments of the above mentioned peptides, containing amino
acids 1-9, 1-10, 1-11, 1-12, or 1-13, are also embodiments of the present
invention. The invention further encompasses pharmaceutically acceptable salts
of the above-described peptides, and N- or C- derivatives of the peptides.

15 This invention also provides pharmaceutical compositions comprising
any of the PTH derivatives described herein and a pharmaceutically acceptable
excipient and/or a pharmaceutically acceptable solution such as saline or a
physiologically buffered solution.

In one aspect, the invention is directed to a pharmaceutical composition
20 comprising the biologically active peptide having the sequence of SEQ. ID. NOs.
1, 5 or any of the above peptides, and a pharmaceutically acceptable carrier.

In another aspect, the invention is directed to a method of making SEQ.
ID. NO. 1, 5 or any of the above peptides, wherein the peptide is synthesized by
solid phase synthesis, liquid phase synthesis, or solution phase synthesis.

25 In another aspect, the invention is directed to a method of making SEQ. ID.
NO. 1, 5 or any of the above peptides, wherein the peptide is protected by FMOC.

This invention also provides a method for treating mammalian conditions
characterized by decreases in bone mass, the method of which comprises
administering to a subject in need thereof an effective bone mass-increasing
30 amount of a biologically active PTH polypeptide derivative. A preferable
embodiment of the invention is drawn to conditions such as osteoporosis,

hyperparathyroidism and hypercalcemia. The types of osteoporosis include, but are not limited to old age osteoporosis and postmenopausal osteoporosis.

In another aspect, the invention is directed to a method for treating mammalian conditions characterized by decreases in bone mass, the method comprising administering to a subject in need thereof an effective bone mass-increasing amount of a biologically active peptide of having the sequence of SEQ. ID. NO. 1 or 5, or any of the above peptides and a pharmaceutically acceptable carrier. Additional preferable embodiments include using an effective amounts of the polypeptide of about 0.01 $\mu\text{g/kg/day}$ to about 1.0 $\mu\text{g/kg/day}$ wherein the polypeptide is administered parenterally, subcutaneously or by nasal insufflation.

This invention also provides a method for determining rates of bone reformation, bone resorption and/or bone remodeling comprising administering to a patient an effective amount of a labeled PTH polypeptide, such as for example, a peptide having the sequence of SEQ. ID. NO. 1, 5 or derivatives thereof and determining the uptake of the peptide into the bone of the patient. The peptide may be labeled with a label selected from the group consisting of: radiolabels, fluorescent labels, bioluminescent labels, or chemiluminescent labels. In a preferable embodiment the radiolabel is ^{125}I or $^{99\text{m}}\text{Tc}$.

Brief Description of the Figures

FIG. 1. α,α - disubstituted amino acid analog structures Aib, Deg, Ac_3c , Ac_5c . Note the restricted ϕ/ψ rotation favors helix formation. (modified from R.Kau. and P. Balaram, *Bioorganic Medicinal Chemistry*, 7 105-117(1999)).

FIG. 2. Effect of single α,α -substituted amino acid substitutions at position 1 or 3 in a PTH(1-14) analog on cAMP-stimulating potency and P1R-binding affinity in HKRK-B28 cells. The peptide $[\text{Ala}^{1,3,12}, \text{Gln}^{10}, \text{Har}^{11}, \text{Trp}^{14}]\text{PTH}(1-14)\text{NH}_2(\text{parent})(\text{SEQ. ID. NO. 13})$ and analogs thereof in which alanine-1 or alanine-3 were substituted with 1-aminocyclopropane-1-carboxylic acid (Ac_3c); 1-aminocyclopentane-1-carboxylic acid (Ac_5c) or α,α -diethylglycine (Deg) were evaluated in HKRK-B28 cells for their capacity to stimulate intracellular cAMP accumulation (A), (C) and the capacity to inhibit binding of ^{125}I -

-10-

[Aib^{1,3},Nle⁸,Gln¹⁰,Har¹¹,Ala¹², Trp¹⁴,Arg¹⁹,Tyr²¹]rPTH(1-21)NH₂ (SEQ. ID. NO. 36) (B), (D). The cAMP responses are expressed as a percent of the maximum response observed in each experiment for the parent peptide, the average of which was 225±21 picomoles of cAMP per well (n=7); the corresponding basal cAMP level was 3.8±0.1 picomoles per well. Peptides and their corresponding symbols are identified in the key.

FIG. 3. Effects of combined α,α-disubstituted amino acid substitutions at positions 1 and 3 in a PTH(1-14) analog on cAMP-stimulating potency and binding affinity in HKRK-B28 cells. The peptide [Ac₃c¹,Aib³,Gln¹⁰,Har¹¹,Ala¹²,Trp¹⁴]PTH(1-14)NH₂ (SEQ. ID. NO. 15) and [Deg^{1,3},Gln¹⁰,Har¹¹,Ala¹²,Trp¹⁴]PTH(1-14)NH₂ (SEQ. ID. NO. 27) were evaluated in HKRK-B28 cells for the capacity to stimulate intracellular cAMP accumulation (A) and the capacity to inhibit the binding of ¹²⁵I-[Aib^{1,3},Nle⁸,Gln¹⁰,Har¹¹,Ala¹², Trp¹⁴,Arg¹⁹,Tyr²¹]rPTH(1-21)NH₂ (SEQ. ID. NO. 36) (B). Also shown is the response observed with the parent peptide [Ala^{1,3,12},Gln¹⁰,Har¹¹,Trp¹⁴]PTH(1-14)NH₂ (SEQ. ID. NO. 13). The cAMP responses are expressed as a percent of the maximum response observed in each experiment for the parent peptide. Peptides and corresponding symbols are identified in the key.

FIG. 4. Phospholipase-C signaling properties of PTH(1-14) analogs in COS-7 cells expressing the hP1R. The capacities of [Ac₃c¹,Aib³,Gln¹⁰,Har¹¹,Ala¹²,Trp¹⁴]PTH(1-14)NH₂ (SEQ. ID. NO. 15), [Aib^{1,3},Gln¹⁰,Har¹¹,Ala¹²,Trp¹⁴]PTH(1-14)NH₂ (SEQ. ID. NO. 3), and [Deg^{1,3},Gln¹⁰,Har¹¹,Ala¹²,Trp¹⁴]PTH(1-14)NH₂ (SEQ. ID. NO. 27) To stimulate formation of 3H-inositol phosphates (IP¹+IP²+IP³) in COS-7 cells transiently transfected with the hP1R, were evaluated. Peptides and corresponding symbols are defined in the key.

FIG. 5. Stimulation of cAMP formation by PTH(1-9) and PTH(1-10) analogs in HKRK-B28 cells. Panel A shows the capacity of varying concentrations of [Ac₃c¹,Aib³,Gln¹⁰]PTH(1-10)NH₂ (SEQ. ID. 22) to stimulate intracellular cAMP accumulation in HKRK-B28 cells. The response to [Ala^{1,3,12},Gln¹⁰,Har¹¹,Trp¹⁴]PTH(1-14)NH₂ (SEQ. ID. NO. 13) is reshown from

-11-

Figure 2A. Panel B shows the cAMP levels in cells treated with buffer alone (basal) or buffer containing either native PTH(1-9)NH₂, [Aib^{1,3}]PTH(1-9)NH₂ (SEQ. ID. NO. 37) or [Ac₅c¹,Aib³]PTH(1-9)NH₂ (SEQ. ID. NO. 34), each at a concentration of 100 μM (*P* versus basal =0.052(*) or <0.0001(a); *P* versus [Aib^{1,3}]PTH(1-9)NH₂ = 0.001(b)(SEQ. ID. NO. 37). Also analyzed were [Ac₅c¹,Aib³,Gln¹⁰]PTH(1-10)NH₂ (SEQ. ID. NO. 22) at 10 μM, and [Ac₅c¹,Aib³,Gln¹⁰,Har¹¹,Ala¹²,Trp¹⁴]PTH(1-14)NH₂ (SEQ. ID.15) at 10 nM, for which 33-fold (*P*=0.0003) and 38-fold (*P*=0.006) increases in cAMP levels, relative to basal, were observed, respectively. Peptides and corresponding symbols are identified in the key.

FIG. 6. cAMP-stimulating capacities of PTH(1-14) and a PTH(1-10) analogs in LdelNt-2 cells. The peptides [Aib^{1,3},Gln¹⁰,Har¹¹,Ala¹²,Trp¹⁴]PTH(1-14)NH₂(SEQ.ID.NO.3), [Ac₅c¹,Aib³,Gln¹⁰,Har¹¹,Ala¹²,Trp¹⁴]PTH(1-14)NH₂(SEQ.ID.NO.15), [Deg^{1,3},Gln¹⁰,Har¹¹,Ala¹²,Trp¹⁴]PTH(1-14)NH₂(SEQ. ID. NO. 27) and [Ac₅c¹,Aib³,Gln¹⁰]PTH(1-10)NH₂ (SEQ. ID. NO. 22) were evaluated for cAMP-stimulating potency in LdelNt-2 cells, which are clonal, LLC-PK1-derived cells that express hP1R-delNt via stable transfection. The data are expressed as a percent of the maximum response observed in each experiment for [Ac₅c¹,Aib³,Gln¹⁰,Har¹¹,Ala¹²,Trp¹⁴]PTH(1-14)NH₂(SEQ. ID. NO.15), which was 125 ± 19 picomoles of cAMP per well; the basal cAMP levels were 1.8±0.3 picomoles per well. The corresponding EC₅₀ values were 6.4±1.6nM ([Aib^{1,3},Gln¹⁰,Har¹¹,Ala¹²,Trp¹⁴]PTH(1-14)NH₂)(SEQ. ID. NO. 3); 6.4±2.3nM([Ac₅c¹,Aib³,Gln¹⁰,Har¹¹,Ala¹²,Trp¹⁴]PTH(1-14)NH₂(SEQ. ID. NO. 15); 1.4±0.5 μM (Deg^{1,3},Gln¹⁰,Har¹¹,Ala¹²,Trp¹⁴]PTH(1-14)NH₂) (SEQ. ID. NO. 27) and 37±22 μM([Ac₅c¹,Aib³,Gln¹⁰]PTH(1-10)NH₂) (SEQ. ID. NO. 22). Peptides and corresponding symbols are identified in the key.

FIG. 7. Stimulation of PLC activity. The PTH(1-14) analog stimulated PLC activity in COS-7 cells transiently expressing hP1R-WT. Peptides and corresponding symbols are identified in the key.

FIG. 8. Agonist activity of PTH analogs *in vivo*. Mice were injected intraperitoneally with vehicle or vehicle containing either [Nle^{8,21},Tyr³⁴]ratPTH(1-

-12-

34)NH₂ (SEQ. ID. NO. 14), or [Ac₃c¹,Aib³,Gln¹⁰,Har¹¹,Ala¹²,Trp¹⁴]humanPTH(1-14)NH₂ (SEQ. ID. NO. 15) and 3 minutes later blood samples were withdrawn and plasma cAMP levels were determined by radioimmunoassay. The effective PTH peptide concentrations were 20x10⁻⁹ mole per kilogram of body-weight for PTH(1-34) analog and 200x10⁻⁹ mole per kilogram of body-weight for PTH(1-14) analog. Each bar represents the mean (±s.e.m.) cAMP value of data obtained from 4 mice.

Detailed Description of the Invention

Definitions

Amino Acid Sequences: The amino acid sequences in this application use either the single letter or three letter designations for the amino acids. These designations are well known to one of skill in the art and can be found in numerous readily available references, such as for example in *Cooper, G.M., The Cell* 1997, ASM Press, Washington, D.C. or Ausubel et al., *Current Protocols in Molecular Biology*, 1994. Where substitutions in a sequence are referred to, for example, as Ser-3 -->Ala or [Ala³]peptide, this means that the serine in the third position from the N-terminal end of the polypeptide is replaced with another amino acid, Alanine in this instance.

Scaffold peptide: [M]PTH(1-14) is defined as [Ala^{1,3,12},Gln¹⁰,Har¹¹,Trp¹⁴]PTH(1-14)amide (SEQ. ID. NO. 13). [M]PTH(1-11) is defined as [Ala^{1,3},Gln¹⁰,Har¹¹]PTH(1-11)NH₂ (SEQ. ID. NO. 16). [Aib^{1,3},M]PTH(1-14) is defined as [Aib^{1,3},Gln¹⁰,Har¹¹,Ala¹²,Trp¹⁴]PTH(1-14)NH₂ (SEQ. ID. NO. 3).

α,α-dialkyl amino acids: "Aib" refers to α-aminoisobutyric acid; "Har" refers to homoarginine; "Nle" refers to norleucine; Ac₃c refers to 1-aminocyclopropane-1-carboxylic acid; Ac₄c refers to 1-amino-cyclobutane-carboxylic acid; Ac₅c refers to 1-aminocyclopentane-1-carboxylic acid; Ac₆c refers to 1-amino-cyclohexane-carboxylic acid; Deg refers to α,α-diethylglycine; IBMX refers to 3-isobutyl-1-methylxanthine; and other amino acids are in either the

-13-

conventional one- or three-letter codes.

Biological Activity of the Protein: This expression refers to any biological activity of the polypeptide. Examples of these activities include, but are not limited to metabolic or physiologic function of compounds of a peptide having the sequence of SEQ. ID. NO. 1 or derivatives thereof, including similar activities or improved activities, or those activities with decreased undesirable side-effects. Also included are antigenic and immunogenic activities of the above-described compounds.

Derivative or Functional Derivative: The term "derivative" or "functional derivative" is intended to include "variants," the "derivatives," or "chemical derivatives" of the PTH molecule. A "variant" of a molecule such as for example, a compound of a peptide having the sequence of SEQ. ID. NO. 1 or derivative thereof is meant to refer to a molecule substantially similar to either the entire molecule, or a fragment thereof. An "analog" of a molecule such as for example, a compound having the sequence of SEQ. ID. NO. 1 or derivative thereof is meant to refer to a non-natural molecule substantially similar to either the peptide having the sequence of SEQ. ID. NO. 1 molecules or fragments thereof.

PTH derivatives contain changes in the polypeptide relative to the native PTH polypeptide of the same size. The sequence of the native human (" h P T H ") P T H (1 - 1 4) p o l y p e p t i d e is SerValSerGluIleGlnLeuMetHisAsnLeuGlyLysHis (part of SEQ. ID. NO. 26), or native rat ("rPTH") PTH (1-14) is AlaValSerGluIleGlnLeuMetHisAsnLeuGly LysHis (SEQ. ID. NO. 17). A molecule is said to be "substantially similar" to another molecule if the sequence of amino acids in both molecules is substantially the same, and if both molecules possess a similar biological activity. Thus, two molecules that possess a similar activity, may be considered variants, derivatives, or analogs as that term is used herein even if one of the molecules contains additional amino acid residues not found in the other, or if the sequence of amino acid residues is not identical. PTH derivatives, however, need not have substantially similar biological activity to the native molecule. In some instances PTH derivatives have substantially different activity than the native PTH. For

-14-

example, a derivative may be either an antagonist or an agonist of the PTH receptor.

As used herein, a molecule is said to be a "chemical derivative" of another molecule when it contains additional chemical moieties not normally a part of the molecule. Such moieties may improve the molecule's solubility, absorption, biological half-life, etc. The moieties may alternatively decrease the toxicity of the molecule, eliminate or attenuate any undesirable side effect of the molecule, etc. Examples of moieties capable of mediating such effects are disclosed in *Remington's Pharmaceutical Sciences* (1980) and will be apparent to those of ordinary skill in the art.

Fragment: A "fragment" of a molecule such as for example, $X_{01}\text{Val}X_{02}\text{GluIleGlnLeuMetHis}X_{03}X_{04}X_{05}X_{06}$ (1-13) (part of SEQ. ID. NO. 1) or derivative thereof is meant to refer to any polypeptide subset of these molecules, including N- or C- derivatives thereof.

Fusion protein: As used herein, a "fusion protein" is a protein comprising compounds such as for example, $X_{01}\text{Val}X_{02}\text{GluIleGlnLeuMetHis}X_{03}X_{04}X_{05}X_{06}X_{07}$ (SEQ. ID. NO. 1), or derivatives thereof, either with or without a "selective cleavage site" linked at its N-terminus, which is in turn linked to an additional amino acid leader polypeptide sequence.

Polypeptide: Polypeptide and peptide are used interchangeably. The term polypeptide refers to any peptide or protein comprising two or more amino acids joined to each other by peptide bonds or modified peptide bonds, *i.e.*, peptide isosteres. "Polypeptide" refers to both short chains, commonly referred to as peptides, oligopeptides or oligomers, and to longer chains, generally referred to as proteins. Polypeptides may contain amino acids other than the 20 gene-encoded amino acids and include amino acid sequences modified either by natural processes, such as post-translational processing, or by chemical modification techniques which are well known in the art. Such modifications are well described in basic texts and in more detailed monographs, as well as in the research literature. Modifications can occur anywhere in a polypeptide, including the peptide backbone, the amino acid side-chains and the amino or carboxyl termini.

It will be appreciated that the same type of modification may be present in the same or varying degrees at several sites in a given polypeptide. Also, a given polypeptide may contain many types of modifications.

Polypeptides may be branched and they may be cyclic, with or without branching. Cyclic, branched and branched cyclic polypeptides may result from post-translational modifications or may be made by synthetic methods. Modifications include acetylation, acylation, ADP-ribosylation, amidation, covalent attachment of flavin, covalent attachment of a heme moiety, covalent attachment of a nucleotide or nucleotide derivative, covalent attachment of a lipid or lipid derivative, covalent attachment of phosphatidylinositol, cross-linking, cyclization, disulfide bond formation, demethylation, formation of covalent cross-links, formation of cystine, formation of pyroglutamate, formylation, gamma-carboxylation, glycosylation, GPI anchor formation, hydroxylation, iodination, methylation, myristoylation, oxidation, proteolytic processing, phosphorylation, prenylation, racemization, selenoylation, sulfation, transfer-RNA mediated addition of amino acids to proteins such as arginylation, and ubiquitination. See, for instance, *Proteins-Structure and Molecular Properties*, 2nd Ed., T. E. Creighton, W. H. Freeman and Company, New York, 1993 and Wold, F., *Posttranslational Protein Modifications: Perspectives and Prospects*, pgs. 1-12 in *Posttranslational Covalent Modification of Proteins*, B. C. Johnson, Ed., Academic Press, New York, 1983; Seifter *et al.*, "Analysis for protein modifications and nonprotein cofactors", *Methods in Enzymol.* 182:626-646 (1990) and Rattan *et al.*, "Protein Synthesis: Posttranslational Modifications and Aging", *Ann NY Acad Sci* 663:48-62 (1992).

PTH Receptor: PTH-1 receptor (P1R) is a class IIG protein coupled receptor, the amino-terminal extracellular (N)domain of which is thought to bind with the C-terminal portion of PTH.

PTH Analogs -- Structural and Functional Properties

α -aminoisobutyric acid (Aib) and α,α -disubstituted amino acids distinct

from Aib were introduced into short N-terminal PTH peptide analogs. The numerous NMR studies of PTH(1-34) analogs, performed in a variety of polar or non-polar solvents, have generally indicated two domains of secondary structure: a stable C-terminal helix extending approximately from Ser-17 to Val-31, and a shorter and less stable amino-terminal helix, extending variably from Ser-3 to Lys-13, the two domain being connected by a bend or turn region (Marx, U.C., *et al.*, *Biochem. Biophys. Res. Commun.* 267:213-220 (2000); Chen, Z., *et al.*, *Biochemistry* 39:12766-12777 (2000); Marx, U.C., *et al.*, *J. Biol. Chem.* 270:15194-15202 (1995); Marx, U.C., *et al.*, *J. Biol. Chem.* 273:4308-4316 (1998); Pellegrini, M., *et al.*, *Biochemistry* 37:12737-12743 (1998); Gronwald, W., *et al.*, *Biol. Chem. Hoppe Seyler* 377:175-186 (1996); Barden, J.A., and Kemp, B.E., *Biochemistry* 32:7126-7132 (1993)). The recent crystallographic study of PTH(1-34) indicated a continuous α -helix extending from Ser-3 to His-32 and containing only a slight 15° bend at the mid-section. However, NMR data indicates that the N-terminal α -helix is relatively weak. Helix-stabilizing modifications, such as the introduction of Aib residues, offer significant benefits in terms of peptide potency, and result in short peptides (≤ 14 amino acids) with activity comparable to PTH(1-34).

Described herein are novel "minimized" variants of PTH that are small enough to be deliverable by simple non-injection methods. The variants of the present invention contain substitutions in the first 14 amino acids of the polypeptide. The new polypeptides correspond to the 1-14, 1-13, 1-12, 1-11, 1-10 and 1-9 amino acid sequence of the mature PTH polypeptide, unless otherwise noted. The shorter variants (\leq PTH1-14) have a molecular weight of less than 2,000 daltons.

As protein products, compounds described herein are amenable to production by the techniques of solution- or solid-phase peptide synthesis. The solid phase peptide synthesis technique, in particular, has been successfully applied in the production of human PTH and can be used for the production of these compounds (for guidance, see Kimura *et al.*, *supra*, and see Fairwell *et al.*, *Biochem.* 22:2691 (1983)). Success with producing human PTH on a relatively

-17-

large scale has been reported by Goud *et al.*, in *J. Bone Min. Res.* 6(8):781 (1991). The synthetic peptide synthesis approach generally entails the use of automated synthesizers and appropriate resin as solid phase, to which is attached the C-terminal amino acid of the desired compounds of peptides having the sequence of SEQ. ID. NO. 1 or derivatives thereof. Extension of the peptide in the N-terminal direction is then achieved by successively coupling a suitably protected form of the next desired amino acid, using either FMOC- or BOC-based chemical protocols typically, until synthesis is complete. Protecting groups are then cleaved from the peptide, usually simultaneously with cleavage of peptide from the resin, and the peptide is then isolated and purified using conventional techniques, such as by reversed phase HPLC using acetonitrile as solvent and tri-fluoroacetic acid as ion-pairing agent. Such procedures are generally described in numerous publications and reference may be made, for example, to Stewart and Young, "Solid Phase Peptide Synthesis," 2nd Edition, Pierce Chemical Company, Rockford, IL (1984). It will be appreciated that the peptide synthesis approach is required for production of such as for example, a peptide having the sequence of SEQ. ID. NO. 1 and derivatives thereof which incorporate amino acids that are not genetically encoded, such as Aib.

Substituents may be attached to the free amine of the N-terminal amino acid of compounds of the present invention standard methods known in the art. For example, alkyl groups, e.g., C₁₋₁₂ alkyl, are attached using reductive alkylation. Hydroxyalkyl groups, e.g. C₁₋₁₂ hydroxyalkyl, are also attached using reductive alkylation wherein the free hydroxy group is protected with a t-butyl ester. Acyl groups, e.g., COE₁, are attached by coupling the free acid, e.g., E₁COOH, to the free amino of the N-terminal amino acid. Additionally, possible chemical modifications of the C-terminal end of the polypeptide are encompassed within the scope of the invention. These modifications may modify binding affinity to the receptor.

Also contemplated within the scope of this invention are those compounds such as for example, a peptide having the sequence of SEQ. ID. NO. 1 and derivatives thereof with altered secondary or tertiary structure, and/or altered

-18-

stability, which still retain biological activity. Such derivatives might be achieved through lactam cyclization, disulfide bonds, or other means known to a person of ordinary skill in the art. A preferable embodiment of the invention is drawn to any of the above recited polypeptides, wherein the polypeptide contains a C-terminal amide.

Utility and Administration of Compounds of the Invention

Compounds of the invention or derivatives thereof have multiple uses. These include, *inter alia*, agonists or antagonists of the PTH receptor, prevention and treatment of a variety of mammalian conditions manifested by loss of bone mass, diagnostic probes, antigens to prepare antibodies for use as diagnostic probes and even as molecular weight markers. Being able to specifically substitute one or more amino acids in the PTH polypeptide permits construction of specific molecular weight polypeptides.

In particular, the compounds of this invention are indicated for the prophylaxis and therapeutic treatment of osteoporosis and osteopenia in humans. Furthermore, the compounds of this invention are indicated for the prophylaxis and therapeutic treatment of other bone diseases. The compounds of this invention are also indicated for the prophylaxis and therapeutic treatment of hypoparathyroidism. Finally, the compounds of this invention are indicated for use as agonists for fracture repair and as antagonists for hypercalcemia.

In general, compounds of the present invention, or salts thereof, are administered in amounts between about 0.01 and 1 $\mu\text{g/kg}$ body weight per day, preferably from about 0.07 to about 0.2 $\mu\text{g/kg}$ body weight per day. For a 50 kg human female subject, the daily dose of biologically active compound is from about 0.5 to about 50 μgs , preferably from about 3.5 to about 10 μgs . In other mammals, such as horses, dogs, and cattle, higher doses may be required. This dosage may be delivered in a conventional pharmaceutical composition by a single administration, by multiple applications, or via controlled release, as needed to achieve the most effective results, preferably one or more times daily by injection.

For example, this dosage may be delivered in a conventional pharmaceutical composition by nasal insufflation or by orally active means. Another method of administration includes subcutaneous injection, at least once daily, for potential anabolic effects on the skeleton.

5 The selection of the exact dose and composition and the most appropriate delivery regimen will be influenced by, *inter alia*, the pharmacological properties of the selected compounds of the invention, the nature and severity of the condition being treated, and the physical condition and mental acuity of the recipient.

10 Representative preferred delivery regimens include, without limitation, oral, parenteral, subcutaneous, transcutaneous, intramuscular and intravenous, rectal, buccal (including sublingual), transdermal, and intranasal insufflation.

 Pharmaceutically acceptable salts retain the desired biological activity of the compounds of the invention without toxic side effects. Examples of such salts
15 are (a) acid addition salts formed with inorganic acids, for example hydrochloric acid, hydrobromic acid, sulfuric acid, phosphoric acid, nitric acid and the like; and salts formed with organic acids such as, for example, acetic acid, oxalic acid, tartaric acid, succinic acid, maleic acid, fumaric acid, gluconic acid, citric acid, malic acid, ascorbic acid, benzoic acid, tannic acid, pamoic acid, alginic acid,
20 polyglutamic acid, naphthalenesulfonic acids, naphthalene disulfonic acids, polygalacturonic acid and the like; (b) base addition salts formed with polyvalent metal cations such as zinc, calcium, bismuth, barium, magnesium, aluminum, copper, cobalt, nickel, cadmium, and the like; or with an organic cation formed from N,N'-dibenzylethylenediamine or ethylenediamine; or (c) combinations of
25 (a) and (b), e.g., a zinc tannate salt and the like. Pharmaceutically acceptable buffers include but are not limited to saline or phosphate buffered saline. Also included in these solutions may be acceptable preservative known to those of skill in the art.

 A further aspect of the present invention relates to pharmaceutical
30 compositions comprising as an active ingredient compounds of the invention or derivatives thereof of the present invention, or pharmaceutically acceptable salt

-20-

thereof, in admixture with a pharmaceutically acceptable, non-toxic carrier. As mentioned above, such compositions may be prepared for parenteral (subcutaneous, transcutaneous, intramuscular or intravenous) administration, particularly in the form of liquid solutions or suspensions; for oral or buccal
5 administration, particularly in the form of tablets or capsules; for rectal, transdermal administration; and for intranasal administration, particularly in the form of powders, nasal drops or aerosols.

The compositions may conveniently be administered in unit dosage form and may be prepared by any of the methods well-known in the pharmaceutical art, for example as described in Remington's Pharmaceutical Sciences, 17th ed., Mack
10 Publishing Company, Easton, Pa., (1985), incorporated herein by reference. Formulations for parenteral administration may contain as excipients sterile water or saline, alkylene glycols such as propylene glycol, polyalkylene glycols such as polyethylene glycol, oils of vegetable origin, hydrogenated naphthalenes and the like. For oral administration, the formulation can be enhanced by the addition of
15 bile salts or acylcarnitines. Formulations for nasal administration may be solid and may contain excipients, for example, lactose or dextran, or may be aqueous or oily solutions for use in the form of nasal drops or metered spray. For buccal administration typical excipients include sugars, calcium stearate, magnesium stearate, pregelatinated starch, and the like.
20

When formulated for one preferred route of administration, nasal administration, the absorption across the nasal mucous membrane may be enhanced by surfactant acids, such as for example, glycocholic acid, cholic acid, taurocholic acid, ethocholic acid, deoxycholic acid, chenodeoxycholic acid,
25 dehydrocholic acid, glycodeoxycholic acid, cyclodextrins and the like in an amount in the range between about 0.2 and 15 weight percent, preferably between about 0.5 and 4 weight percent, most preferably about 2 weight percent.

Delivery of the compounds of the present invention to the subject over prolonged periods of time, for example, for periods of one week to one year, may
30 be accomplished by a single administration of a controlled release system containing sufficient active ingredient for the desired release period. Various

-21-

controlled release systems, such as monolithic or reservoir-type microcapsules, depot implants, osmotic pumps, vesicles, micelles, liposomes, transdermal patches, iontophoretic devices and alternative injectable dosage forms may be utilized for this purpose. Localization at the site to which delivery of the active ingredient is desired is an additional feature of some controlled release devices, which may prove beneficial in the treatment of certain disorders.

One form of controlled release formulation contains the polypeptide or its salt dispersed or encapsulated in a slowly degrading, non-toxic, non-antigenic polymer such as copoly(lactic/glycolic) acid, as described in the pioneering work of Kent, Lewis, Sanders, and Tice, U.S. Pat. No. 4,675,189. The compounds or, preferably, their relatively insoluble salts, may also be formulated in cholesterol or other lipid matrix pellets, or silastomer matrix implants. Additional slow release, depot implant or injectable formulations will be apparent to the skilled artisan. See, for example, Sustained and Controlled Release Drug Delivery Systems, J. R. Robinson ed., Marcel Dekker, Inc., New York, 1978, and R. W. Baker, Controlled Release of Biologically Active Agents, John Wiley & Sons, New York, 1987.

Like PTH, the PTH variants may be administered in combination with other agents useful in treating a given clinical condition. When treating osteoporosis and other bone-related disorders for example, the PTH variants may be administered in conjunction with a dietary calcium supplement or with a vitamin D analog (*see* U.S. Pat. No. 4,698,328). Alternatively, the PTH variant may be administered, preferably using a cyclic therapeutic regimen, in combination with bisphosphonates, as described for example in U.S. Pat. No. 4,761,406, or in combination with one or more bone therapeutic agents such as, without limitation, calcitonin and estrogen.

PTH Analog Receptor-Signaling Activities

A crucial step in the expression of hormonal action is the interaction of hormones with receptors on the plasma membrane surface of target cells. The formation of hormone-receptor complexes allows the transduction of extracellular

-22-

signals into the cell to elicit a variety of biological responses.

Polypeptides described herein can be screened for their agonistic or antagonistic properties using the cAMP accumulation assay. Cells expressing PTH-1 receptor on the cell surface are incubated with native PTH(1-84) (SEQ. ID. NO. 18) for 5-60 minutes at 37°C., in the presence of 2 mM IBMX (3-isobutyl-1-methyl-xanthine, Sigma, St. Louis, MO). Cyclic AMP accumulation is measured by specific radio-immunoassay. A compound that competes with native PTH(1-84) or PTH(1-34) (SEQ. ID. NO.19) for binding to the PTH-1 receptor, and that inhibits the effect of native PTH(1-84) or PTH(1-34) on cAMP accumulation, is considered a competitive antagonist. Such a compound would be useful for treating hypercalcemia.

Conversely, a PTH analog described herein or a derivative thereof that does not compete with native PTH(1-84) or PTH(1-34) for binding to the PTH-1 receptor, but which still prevents native PTH(1-84) or PTH(1-34) activation of cAMP accumulation (presumably by blocking the receptor activation site) is considered a non-competitive antagonist. Such a compound would also be useful for treating hypercalcemia.

The compounds described herein that compete with native PTH(1-84) or PTH(1-34) for binding to the PTH-1 receptor, and which stimulates cAMP accumulation in the presence or absence of native PTH(1-84) or PTH(1-34) are competitive agonists. A compound that does not compete with native PTH(1-84) or PTH(1-34) for binding to the PTH-1 receptor but which is still capable of stimulating cAMP accumulation in the presence or absence of native PTH(1-84) or PTH(1-34), or which stimulates a higher cAMP accumulation than that observed by a compound of the invention or a derivative thereof alone, would be considered a non-competitive agonist.

Therapeutic Uses of PTH Analogs

Some forms of hypercalcemia and hypocalcemia are related to the interaction between PTH and PTHrP and the PTH-1 and receptors. Hypercalcemia

is a condition in which there is an abnormal elevation in serum calcium level; it is often associated with other diseases, including hyperparathyroidism, osteoporosis, carcinomas of the breast, lung and prostate, epidermoid cancers of the head and neck and of the esophagus, multiple myeloma, and hypernephroma.

5 Hypocalcemia, a condition in which the serum calcium level is abnormally low, may result from a deficiency of effective PTH, *e.g.*, following thyroid surgery.

By "agonist" is intended a ligand capable of enhancing or potentiating a cellular response mediated by the PTH-1 receptor. By "antagonist" is intended a ligand capable of inhibiting a cellular response mediated by the PTH-1 receptor.

10 Whether any candidate "agonist" or "antagonist" of the present invention can enhance or inhibit such a cellular response can be determined using art-known protein ligand/receptor cellular response or binding assays, including those described elsewhere in this application.

In accordance with yet a further aspect of the invention, there is provided

15 a method for treating a medical disorder that results from altered or excessive action of the PTH-1 receptor, comprising administering to a patient therapeutically effective amount of a compound of the invention or a derivative thereof sufficient to inhibit activation of the PTH-1 receptor of said patient.

In this embodiment, a patient who is suspected of having a disorder

20 resulting from altered action of the PTH-1 receptor can be treated using compounds of the invention or derivatives thereof of the invention which are selective antagonists of the PTH-1 receptor. Such antagonists include compounds of the invention or derivatives thereof of the invention which have been determined (by the assays described herein) to interfere with PTH-1 receptor-

25 mediated cell activation or other derivatives having similar properties.

To administer the antagonist, the appropriate compound of the invention or a derivative thereof is used in the manufacture of a medicament, generally by being formulated in an appropriate carrier or excipient such as, *e.g.*, physiological saline, and preferably administered intravenously, intramuscularly,

30 subcutaneously, orally, or intranasally, at a dosage that provides adequate inhibition of a compound of the invention or a derivative thereof binding to the

-24-

PTH-1 receptor. Typical dosage would be 1 ng to 10 mg of the peptide per kg body weight per day.

In accordance with yet a further aspect of the invention, there is provided a method for treating osteoporosis, comprising administering to a patient a therapeutically effective amount of a compound of the invention or a derivative thereof, sufficient to activate the PTH-1 receptor of said patient. Similar dosages and administration as described above for the PTH/PTHrP antagonist, can be used for administration of a PTH/PTHrP agonist, *e.g.*, for treatment of conditions such as osteoporosis, other metabolic bone disorders, and hypoparathyroidism and related disorders.

It will be appreciated by those skilled in the art that the invention can be performed within a wide range of equivalent parameters of composition, concentration, modes of administration, and conditions without departing from the spirit or scope of the invention or any embodiment thereof.

Having now fully described the invention, the same will be more readily understood by reference to specific examples which are provided by way of illustration, and are not intended to be limiting of the invention, unless herein specified.

Examples

The following protocols and experimental details are referenced in the examples that follow.

Example 1

Materials and Methods

Peptides. The amino acid sequences of the peptides used in the study were all derived from human or rat PTH sequences and contain a free amino-terminus and an amidated C-terminus. The parent peptide used as a starting scaffold was [M]PTH(1-14), which is defined as [Ala^{1,3,12}, Gln¹⁰, Har¹¹, Trp¹⁴]PTH(1-14)NH₂ (SEQ. ID. NO. 13). Peptides were prepared on automated peptide synthesizers

(model 430A PE, Applied Biosystems, Foster City, CA, or Model 396 MBS Advanced Chem Tect, Louisville, KY) using Fmoc main-chain protecting group chemistry, HBTU/HOBt/DIEA (1:1:2 molar ratio) for coupling reactions, and TFA-mediated cleavage/sidechain-deprotection (MGH Biopolymer Synthesis Facility, Boston, MA). All peptides were desalted by adsorption on a C18-containing cartridge, and purified further by HPLC. The dry peptide powders were reconstituted in 10 mM acetic acid and stored at -80°C. The purity, identity, and stock concentration for each peptide was secured by analytical HPLC, Matrix-assisted laser desorption/ionization (MALDI) mass spectrometry and amino acid analysis. Radiolabeling of [M]PTH(1-21) (SEQ.ID.NO.20) and [Aib^{1,3},M]PTH(1-21) (SEQ.ID.NO.21) was performed using ¹²⁵I-Na (2,200 Ci/mmol, NEN) and chloramine-T; the resultant radioligands were purified by HPLC.

Cell Culture. The cell line HKRK-B28 (Takasu, H., *et al.*, *J. Bone Miner. Res.* 14:11-20 (1999)) was derived from the porcine kidney cell line, LLC-PK₁ by stable transfection with plasmid DNA encoding a recombinant P1R chimera comprised of the opossum P1R from the N-terminus to the mid region of TM3 and the rat P1R from the mid region of TM3 to the C-terminus. The surface density of the P1R in these cells is ~280,000 receptors per cell. The clonal cell line LdelNt-2 was derived from LLC-PK₁ cells via stable transfection with a plasmid encoding P1R-delNt, a recombinant human PTH-1 receptor construct in which most of the amino-terminal extracellular domain is deleted. These cells, as well as COS-7 cells and SaOS-2-B10 cells, were cultured at 27°C in T-75 flasks (75 mm²) in Dulbecco's modified Eagle's medium (DMEM) supplemented with fetal bovine serum (10%), penicillin G (20 units/ml), streptomycin sulfate (20 µg/ml) and amphotericin B (0.05 µg/ml) in a humidified atmosphere containing 5% CO₂. Stock solutions of EGTA/trypsin and antibiotics were from GIBCO; fetal bovine serum was from Hyclone Laboratories (Logan, UT). COS-7 cells sub-cultured in 24-well plates were transfected with plasmid DNA (200 ng per well) encoding the wild-type human P1R or truncated human P1R deleted for residues (24-181) (Shimizu, M., *et al.*, *J. Biol. Chem.* 275:21836-21843 (2000)) that was purified by cesium chloride/ethidium bromide density gradient centrifugation, and FuGENE 6

-26-

transfection reagent (Roche Indianapolis IN) according to the manufacturer's recommended procedure. All cells, in 24-well plates, were treated with fresh media and shifted to 33°C for 12 to 24 h prior to assay.

5 cAMP Stimulation. Stimulation of cells with peptide analogs was performed in 24-well plates. Cells were rinsed with 0.5 mL of binding buffer (50 mM Tris-HCl, 100 mM NaCl, 5 mM KCl, 2 mM CaCl₂, 5% heat-inactivated horse serum, 0.5% fetal bovine serum, adjusted to pH 7.5 with HCl) and treated with 200 μL of cAMP assay buffer (Delbecco's modified Eagle's medium containing 2 mM 3-isobutyl-1-methylxanthine, 1 mg/mL bovine serum albumin, 35 mM Hepes-
10 NaOH, pH 7.4) and 100 μL of binding buffer containing varying amounts of peptide analog (final volume = 300 μL). The medium was removed after incubation for 30 to 60 minutes at room temperature, and the cells were frozen on dry ice, lysed with 0.5 mL 50 mM HCl, and refrozen (~80°C). The cAMP content of the diluted lysate was determined by radioimmunoassay. The EC₅₀ response
15 values were calculated using nonlinear regression (see below).

Fmoc. (Fluorenylmethoxycarbonyl group) A group used for linkage to amino groups for the purpose either of forming fluorescent amino-acid derivatives that can readily be detected after column chromatography, or to protect the amino groups of amino acids or nucleotides while other functional groups are undergoing
20 reaction. Reagents useful for introducing the group are 9-fluorenylmethyl chloroformate and 9-fluorenyl-methyl succinimidyl carbonate.

Competition Binding. Binding reactions were performed with HKRK-B28 cells or in COS-7 cells in 24-well plates. The cells were rinsed with 0.5 mL of binding buffer, and then treated successively with 100 μL binding buffer, 100 μL
25 of binding buffer containing various amounts of unlabeled competitor ligand, and 100 μL of binding buffer containing ca. 100,000 cpm of ¹²⁵I-[Aib^{1,3},Nle⁸,Gln¹⁰,Har¹¹,Ala¹²,Trp¹⁴, Arg¹⁹,Tyr²¹]rPTH(1-21) (SEQ. ID. NO. 36) (ca. 26 fmol; final volume = 300 μL). Incubations were 4 to 6 h at 4°C, at which time near equilibrium conditions were attained. Cells were then placed on ice, the
30 binding medium was removed, and the monolayer was rinsed three times with 0.5 mL of cold binding buffer. The cells were subsequently lysed with 0.5 mL 5N

-27-

NaOH and counted for radioactivity. For each tracer and in each experiment, the non-specific binding was determined as the radioactivity that bound in the presence of the same unlabeled peptide at a concentration of 1 μ M, and was ~1% of total radioactivity added for each tracer. The maximum specific binding (B_0) was the total radioactivity bound in the absence of competing ligand, corrected for nonspecific binding, and for each tracer, ranged from 8% to 20% of the total radioactivity added. Nonlinear regression was used to calculate binding IC_{50} values (see below). Scatchard transformations of homologous competition binding data derived from studies with 26 fmol of 125 I-[Aib^{1,3},Nle⁸,Gln¹⁰,Har¹¹,Ala¹²,Trp¹⁴, Arg¹⁹,Tyr²¹]rPTH(1-21) (SEQ. ID. NO. 36) were employed for estimations of apparent equilibrium dissociation constant (K_{Dapp} s) and total number of ligand binding sites (B_{max}), assuming a single class of binding sites and equal affinities of the iodinated and non iodinated ligand.

Stimulation of Inositol Phosphate Production. COS-7 cells transfected as above with P1R-WT were treated with serum-free, inositol-free DMEM containing 0.1% bovine serum albumin and [3 H]myo-inositol (NEN, Boston, MA) (2 μ Ci/mL) for 16 h prior to assay. At the time of the assay, the cells were rinsed with binding buffer containing LiCl (30 mM) and treated with the same buffer with or without a PTH analog. The cells were then incubated at 37°C for 40 min, after which the buffer was removed and replaced by 0.5 mL of ice cold 5% trichloroacetic acid solution. After 3 h on ice, the lysate was collected and extracted twice with ethyl ether. The lysate was then applied to an ion exchange column (0.5 mL resin bed) and the total inositol phosphates were eluted as described previously (Berridge, M.J., *et al.*, *Biochem. J.* 212:473-482 (1983)), and counted in liquid scintillation cocktail.

Inhibition of Chondrocyte Differentiation in Embryonic Mouse Metatarsals. Metatarsals from embryonic day (E) 15.5 mouse embryos were excised and cultured in a 37°C humidified incubator (5% CO₂) in serum-free α MEM media in 24 well plates. Sixteen hours later, a PTH analog or vehicle was added, and the samples were incubated for an additional 48 h in 37°C with peptide or vehicle added again at the 24 h time point. At the end of the 64 h incubation

period, the samples were fixed with 10% formalin/phosphate-buffered saline, then directly visualized on a dissecting microscope using white light. Sections were processed for in-situ hybridization analysis using ³⁵S-labeled riboprobes specific for collagen type X mRNA, a developmental marker gene expressed only in hypertrophic chondrocytes of the growth plate.

Circular Dichroism. Circular Dichroism spectra were recorded on a Jasco model 710 spectropolarimeter; peptides were analyzed at a concentration of 20 μ M in 50 mM sodium phosphate buffer pH 7.4, or the same buffer containing 2,2,2-trifluoroethanol at 20% (v/v). Spectroscopic scans were performed at 20°C and at wavelengths between 185 and 255 nm, with data recorded at each 1 nm interval. The spectral bandwidth was 1.5 nm and 8 scans were accumulated and averaged for each sample. At each wavelength, the mean residue ellipticity [$\theta \times 100 / l \times C \times n$]; where θ is the raw ellipticity value (in dimensions of millidegree), l is the sample path length, C is the molar peptide concentration, and n is the number of residues in the peptide (Bowen, W.P., and Jerman, J.C., *Trends in Pharmacol. Sci.* 16: 413-417 (1995)). The helical content of each peptide was estimated by dividing $[\theta]$ observed at 222 nm for that peptide by -28,100, which is the reported $[\theta]_{222}$ obs for a model helical decapeptide (Bowen, W.P., and Jerman, J.C., *Trends in Pharmacol. Sci.* 16: 413-417 (1995)).

Data Calculation. Calculations were performed using Microsoft® Excel. Nonlinear regression analyses of binding and cAMP dose-response data were performed using the four-parameter equation: $y_p = \text{Min} + [(\text{Max} - \text{Min}) / (1 + (\text{IC}_{50}/x)^{\text{slope}})]$. The Excel Solver function was utilized for parameter optimization, as described previously (Carter, P.H., et al., *Endocrinology* 140: 4972-4981 (1999); Bowen, W.P., and Jerman, J.C., *Trends in Pharmacol. Sci.* 16: 413-417 (1995)). Differences between paired data sets were statistically evaluated using a one-tailed Student's t-test, assuming unequal variances for the two sets.

Example 2

P1-R Binding Affinity of Analogs Containing α,α -disubstituted Amino Acids

5 The effects of introducing α,α -disubstituted amino acids distinct from Aib at positions 1 and/or 3 of [M]PTH(1-14) (M=Ala^{1,3,12}, Gln¹⁰, Har¹¹, Trp¹⁴) (SEQ. ID. NO. 13) were analyzed. Six amino acids were chosen: α -Amino-isobutyric acid (Aib), α,α -diethylglycine (Deg); 1-aminocyclopropane-1-carboxylic acid (Ac₃c); 1-amino-cyclobutane-carboxylic acid (Ac₄c), 1-aminocyclopentane-1-carboxylic acid (Ac₅c), and 1-amino-cyclohexane-carboxylic acid (Ac₆c) (some of which are
10 shown in Fig. 1).

In competition binding assays performed in an LLC-PK1-derived cell line (B28) that stably expresses the hPTH receptor using ¹²⁵I-[Aib^{1,3},Nle⁸,Gln¹⁰,Har¹¹,Ala¹²,Trp¹⁴, Arg¹⁹,Tyr²¹]rPTH(1-21) (SEQ. ID. NO. 36) tracer, the affinities of the Deg³- and Ac₅c³-containing analog were comparable to
15 that of [M]PTH(1-14) (IC₅₀s ~3 μ M). The affinity of Ac₃c³-containing analog was 14-fold lower than that of [M]PTH(1-14), and the affinities of the Deg-, Ac₃c- and Ac₅c-substituted peptides were 3-50 fold higher than that of [M]PTH(1-14) (IC₅₀=0.6 μ M).

The analog [Ac₃c¹,Aib³,M]PTH(1-14) (SEQ. ID. NO. 15) exhibited one of
20 the highest binding affinities (IC₅₀=100 nM) and one of the highest cAMP potencies {EC₅₀=0.9 nM, compared to 200 nM for [M]PTH(1-14)} of any PTH(1-14) analogs studied to date. The Ac₃c¹ modification improved affinity 50-fold relative to the parent [M]PTH(1-14) analog {[Ala^{1,3,12},Gln¹⁰,Har¹¹, Trp¹⁴]humanPTH(1-14)-NH₂} (SEQ. ID. NO. 13), and it improved cAMP potency
25 ~35 fold. A cyclopentane ring at position -1 therefore enables a more favorable interaction with the P1R than do the two C α -methyl of Aib. That each of these structurally distinct amino acids at position 1 improves affinity/potency suggests that their enhancing effects are not due to their specific side chain topologies, but rather to their effects on backbone conformation, that is stabilization of α -helix
30 structure. Deg³ substitution showed marginal improvement of affinity by around 1.5 fold, but markedly diminished signaling potency by ~40 fold.

-30-

Example 3

Ac₃c Peptide Agonist Activity

The peptide [Ac₃c¹, Aib³, M]PTH(1-14)(SEQ. ID. NO. 15) was ~ 2-fold more potent than [Aib^{1,3},M]PTH(1-14) (SEQ. ID. NO. 3) for inhibiting ¹²⁵I-[Aib^{1,3},Nle⁸,Gln¹⁰,Har¹¹,Ala¹²,Trp¹⁴, Arg¹⁹,Tyr²¹]rPTH(1-21) (SEQ. ID. NO. 36) binding to B28 cells, and for stimulating adenylyl cyclase, as well as for stimulating phospholipase C (Fig. 2). This peptide is thus one of the most potent PTH(1-14) analog identified so far. Combining Deg¹ and Deg³ yielded a peptide that bound with adequate affinity (Fig. 1A) but was a true partial agonist for cAMP formation (Fig. 1B).

The peptide [Ac₃c¹, Aib³, Gln¹⁰]PTH(1-10) (SEQ. ID. NO. 22) was a full cAMP agonist, albeit with micromolar potency at 10⁻⁴M, [Ac₃c¹, Aib³]PTH (1-9) (SEQ. ID. NO. 31) exhibited clear agonist activity making it the shortest PTH analog peptide with reliable cAMP agonist activity.

Example 4

Ac₃c Peptide Antagonist Activity

Ligands that function as antagonists can be useful for treating hyperparathyroidism. Analogs of [Ac₃c¹, Aib³, M]PTH(1-14)(parent) (SEQ. ID. NO. 15) containing N-terminal modifications predicted to dissociate signaling and binding affinity (e.g. desamino-1, Trp-2, Bpa-2, Arg-2, Deg-1,3). Most of the substitutions moderately reduced P1R binding affinity and strongly reduced cAMP-stimulating potency.

The Trp² analog [Ac₃c¹,Trp²,Aib³, M]PTH(1-14) (SEQ. ID. NO. 23) at 10⁻⁵M inhibits the agonist activity of [Aib^{1,3},M]PTH(1-14) (SEQ. ID. NO. 3) at 10⁻⁹M by ~50%.

Example 5

Single Substitutions at Positions 1 and 3 in PTH (1-14)

-31-

Single substitutions of di-alkyl amino acids were introduced at positions 1 and 3 in the parent scaffold peptide [Ala^{1,3,12},Gln¹⁰,Har¹¹,Trp¹⁴]PTH(1-14)NH₂ (SEQ.ID.NO.13). Schematic structures of the amino acids utilized at positions 1 and 3 in these studies are shown in Figure 1, and the peptide sequences are presented in Table 1. The parent peptide [Ala^{1,3,12},Gln¹⁰,Har¹¹,Trp¹⁴]PTH(1-14)NH₂ (SEQ. ID. NO. 13), which contains alanine at positions 1 and 3, stimulated cAMP accumulation in HKRK-B28 cells with a potency (EC₅₀) value of 220±80nM, and it inhibited the binding of ¹²⁵I-[Aib^{1,3},Nle⁸,Gln¹⁰,Har¹¹,Ala¹²,Trp¹⁴,Arg¹⁹,Tyr²¹]rPTH(1-21) (SEQ. ID. NO. 36) tracer radioligand to these cells with an apparent affinity (IC₅₀) value of 27±3μM. Relative to the parent analog, the analogs substituted at position 1 were ~two-fold (Ac₃c, *P*=0.1), 11-fold (Deg, *P*=0.02) or 61-fold (Ac₅c, *P*=0.02) more potent for stimulating cAMP accumulation, and these increases in potency were accompanied by commensurate effects on apparent binding affinity. At position 3, substitution with either cycloalkane amino acid, Ac₃c or Ac₅c, increased cAMP-stimulating potency modestly (<two-fold), whereas substitution with the linear amino acid, Deg, diminished potency approximately 10-fold; Ac₅c-3 and Deg-3 had little or no effect on binding affinity, whereas Ac₃c-3 reduced affinity approximately 10-fold.

Example 6

Combined Substitutions at Positions 1 and 3 in PTH(1-14) analogs

Di-alkyl amino acids were introduced at both positions 1 and 3 of the parent scaffold peptide [Ala^{1,3,12},Gln¹⁰,Har¹¹,Trp¹⁴]PTH(1-14)NH₂ (SEQ.ID.NO.13). Introducing Ac₃c at positions 1 and 3 yielded the analog [Ac₃c^{1,3},Gln¹⁰,Har¹¹,Ala¹²,Trp¹⁴]PTH(1-14)NH₂ (SEQ. ID. NO. 28), which was comparable to the parent peptide in terms of binding affinity and signaling potency. Introducing Ac₅c at these positions yielded the analog [Ac₅c^{1,3},Gln¹⁰,Har¹¹,Ala¹²,Trp¹⁴]PTH(1-14)NH₂ (SEQ. ID. NO. 29), which bound to the P1R with 20-fold higher affinity than did the parent peptide (*P*=0.001) and was 30-fold more potent for cAMP signaling (*P*=0.02; Table 2). Combining the

-32-

Deg substitutions at positions 1 and 3 yielded [Deg^{1,3},Gln¹⁰,Har¹¹,Ala¹²,Trp¹⁴]PTH(1-14)NH₂ (SEQ. ID. NO. 27), which bound to the P1R with an affinity 50-fold higher than that of the parent ($P=0.001$) but elicited only a partial agonist response which attained a maximum that was only 45% of that attained by the parent peptide.

Example 7

PTH(1-9) and PTH(1-10) Analogs

Minimum-length active analogs were developed to be used to functionally probe the PTH•PTH receptor interaction mechanism. Native N-terminal PTH analogs that are shorter than PTH(1-14) are inactive in cell-based assays, measurable cAMP responses can be detected in P1R-transfected cells for the analog [Aib^{1,3},Gln¹⁰]PTH(1-10)NH₂ (SEQ. ID. NO.35). [Ac₃c¹,Aib³,Gln¹⁰]PTH(1-10)NH₂ (SEQ. ID. NO. 22) and [Ac₃c¹,Aib³]PTH(1-9)NH₂ (SEQ. ID. NO. 31) were prepared and their activity was assessed in HKRK-B28 cells. When tested at concentrations as high as 1×10^{-4} M, neither peptide inhibited the binding of P(1-21) tracer radioligand. Each analog induced a clear cAMP response in these cells. From dose-response analysis of [Ac₃c¹,Aib³,Gln¹⁰]PTH(1-10)NH₂ (SEQ. ID. NO. 22), EC₅₀ value was estimated at approximately 3 μ M. The PTH(1-19) analog was assayed at a single high concentration (1×10^{-4} M) and found to induce a six-fold increase in cAMP levels, relative to the basal levels ($P=0.001$); at the same concentrations, [Aib^{1,3},Gln¹⁰]PTH(1-10)NH₂ (SEQ. ID. NO. 35) induced a two-fold increase in cAMP accumulation ($P=0.52$) and native PTH(1-9) (SEQ. ID. NO. 33) was inactive.

Example 8

Analog Activity in P1R-delNT cells

Agonist activities of selected analogs in cells expressing P1R-delNt were examined. This PTH-1 receptor construct lacks most of the N domain, but nevertheless mediates full agonist responses to N-terminally modified PTH

-33-

analogues, and thus enables the relative roles of the receptor's N and J domains to the ligand interaction process to be evaluated. The clonal cell line LdelNT-2 was used, which was derived from LLC-PK₁ cells by stable transfection with a plasmid encoding the human P1R-delNt construct. In these cells, [Ac₅c¹, Aib³, Gln¹⁰, Har¹¹, Ala¹², Trp¹⁴]PTH(1-14)NH₂ (SEQ. ID. NO. 15) induced a 70-fold increase in cAMP accumulation with high potency (EC₅₀=6.4±2.3nM); [Ac₅c¹, Aib³, Gln¹⁰]PTH(1-10)NH₂ (SEQ. ID. NO. 22) exhibited weaker potency (EC₅₀~40μM) and [Deg¹³, Gln¹⁰, Har¹¹, Ala¹², Trp¹⁴]PTH(1-14)NH₂ (SEQ. ID. NO. 27) exhibited partial agonist behavior (E_{max}=23-fold over basal; EC₅₀=1.4±0.5μM).

Table 1: PTH Peptide Sequences

		1	5	10	14
	PTH(1-14)NH ₂	Ser-V-Ser-E-I-Q-L-M-H-N-Leu-G-K-H(SEQ. ID. NO. 26)			
	[M]PTH(1-14) (Parent)	A-V-A-E-I-Q-L-M-H-Q-Har-A-K-W (SEQ. ID. NO. 13)			
	[Deg ¹ ,M]PTH(1-14)	Deg-V-A-E-I-Q-L-M-H-Q-Har-A-K-W(SEQ. ID. NO.24)			
5	[Ac ₃ c ¹ ,M]PTH(1-14)	Ac ₃ c-V-A-E-I-Q-L-M-H-Q-Har-A-K-W(SEQ. ID. NO. 25)			
	[Ac ₃ c ¹ ,M]PTH(1-14)	Ac ₃ c-V-A-E-I-Q-L-M-H-Q-Har-A-K-W(SEQ. ID. NO. 4)			
	[Deg ^{1,3} ,M]PTH(1-14)	Deg-V-Deg-E-I-Q-L-M-H-Q-Har-A-K-W(SEQ. ID. NO. 27)			
	[Ac ₃ c ^{1,3} ,M]PTH(1-14)	Ac ₃ c-V-Ac ₃ c-E-I-Q-L-M-H-Q-Har-A-K-W (SEQ. ID. NO. 28)			
	[Ac ₃ c ^{1,3} ,M]PTH(1-14)	Ac ₃ c-V-Ac ₃ c-E-I-Q-L-M-H-Q-Har-A-K-W (SEQ. ID. NO. 29)			
	[Aib ^{1,3} ,M]PTH(1-14)	Aib-V-Aib-E-I-Q-L-M-H-Q-Har-A-K-W (SEQ.ID. NO.3)			
10	[Ac ₃ c ¹ ,Aib ³ ,M]PTH(1-14)	Ac ₃ c-V-Aib-E-I-Q-L-M-H-Q-Har-A-K-W (SEQ.ID. NO. 15)			
	[Ac ₃ c ¹ ,Aib ³ ,M]PTH(1-10)	Ac ₃ c-V-Aib-E-I-Q-L-M-H-Q (SEQ. ID. NO. 22)			
	[Ac ₃ c ¹ ,Aib ³ ,M]PTH(1-9)	Ac ₃ c-V-Aib-E-I-Q-L-M-H (SEQ. ID. NO. 31)			
	[Ac ₄ c ¹ , Aib ³ ,M]PTH(1-14)	Ac ₄ c-V-Aib-E-I-Q-L-M-H-Q-Har-A-K-W (SEQ. ID. NO. 7)			
	[Ac ₆ c ¹ , Aib ³ ,M]PTH(1-14)	Ac ₆ c-V-Aib-E-I-Q-L-M-H-Q-Har-A-K-W (SEQ. ID. NO. 8)			
15	[Ac ₃ c ¹ , Ac ₄ c ³ ,M]PTH(1-14)	Ac ₃ c-V-Ac ₄ c-E-I-Q-L-M-H-Q-Har-A-K-W (SEQ. ID. NO.9)			
	[Ac ₃ c ¹ , Ac ₆ c ³ ,M]PTH(1-14)	Ac ₃ c-V-Ac ₆ c-E-I-Q-L-M-H-Q-Har-A-K-W (SEQ. ID. NO. 10)			
	[Ac ₄ c ¹ , Ac ₄ c ³ ,M]PTH(1-14)	Ac ₄ c-V-Ac ₄ c-E-I-Q-L-M-H-Q-Har-A-K-W (SEQ. ID. NO. 11)			
	[Ac ₆ c ¹ , Ac ₃ c ³ ,M]PTH(1-14)	Ac ₆ c-V-Ac ₆ c-E-I-Q-L-M-H-Q-Har-A-K-W (SEQ. ID. NO. 12)			
	Tracer Radioligands:	Aib-V-Aib-E-I-Q-L-Nle-H-Q-Har-A-K-W-L-A-S-V-R-R-Y*			
20	¹²⁵ I-[M*]PTH(1-21)	(SEQ. ID. NO. 36)			
		* indicates the iodinated tyrosine			

Having now fully described the present invention in some detail by way of illustration and example for purposes of clarity of understanding, it will be obvious to one of ordinary skill in the art that the same can be performed by modifying or changing the invention with a wide and equivalent range of conditions, formulations and other parameters thereof, and that such modifications or changes are intended to be encompassed within the scope of the appended claims.

All publications, patents and patent applications mentioned herein above are herein incorporated in their entirety and by reference to the same extent as if

each individual publication, patent or patent application was specifically and individually indicated to be incorporated by reference.

-36-

WHAT IS CLAIMED IS:

1. A biologically active peptide consisting essentially of the formula selected from:

- 5 (a) $X_{01}\text{Val}X_{02}\text{GluIleGlnLeuMetHis}X_{03}X_{04}X_{05}X_{06}X_{07}$ (SEQ. ID. NO.1);
 (b) peptides containing amino acids 1-9, 1-10, 1-11, 1-12, or 1-13;
 (c) pharmaceutically acceptable salts of (a) or (b); or
 (d) N – or C-derivatives of (a), (b) or (c);

wherein:

- 10 X_{01} is an α -helix-stabilizing residue, Gly, Ser or Ala;
 X_{02} is an α -helix-stabilizing residue, Ala or Ser;
 X_{03} is Ala, Gln or Asn;
 X_{04} is Arg, Har or Leu;
 X_{05} is an α -helix stabilizing residue, Ala or Gly;
 X_{06} is an α -helix stabilizing residue or Lys;
 15 X_{07} is an α -helix stabilizing residue, Trp or His;

wherein at least one of X_{01} , X_{02} , X_{05} , X_{06} or X_{07} is an α -helix stabilizing residue, and wherein at least one of said α -helix stabilizing residues is Aib, Ac_3c , Ac_4c , Ac_5c , Ac_6c , or Deg.

2. The peptide of claim 1, wherein said peptide is selected from:

- 20 (a) $\text{AlaValDegGluIleGlnLeuMetHisGlnHarAlaLysTrp}$ (SEQ. ID. NO. 37);
 (b) peptides containing amino acids 1-9, 1-10, 1-11, 1-12 or 1-13;
 (c) pharmaceutically acceptable salts of (a) or (b); or
 (d) N - or C - derivatives of (a), (b) or (c).

25 3. The peptide of claim 1, wherein said peptide is selected from:

- (a) $\text{AlaValAc}_3\text{cGluIleGlnLeuMetHisGlnHarAlaLysTrp}$ (SEQ. ID. NO. 38);
 (b) peptides containing amino acids 1-9, 1-10, 1-11, 1-12 or 1-13;
 (c) pharmaceutically acceptable salts of (a) or (b); or

-37-

- (d) N - or C - derivatives of (a), (b) or (c).
4. The peptide of claim 1, wherein said peptide is selected from:
- (a) AlaValAc₃cGluIleGlnLeuMetHisGlnHarAlaLysTrp (SEQ. ID. NO. 39);
- 5 (b) peptides containing amino acids 1-9, 1-10, 1-11, 1-12 or 1-13;
- (c) pharmaceutically acceptable salts of (a) or (b); or
- (d) N - or C - derivatives of (a), (b) or (c).
5. The peptide of claim 1, wherein said peptide is selected from:
- (a) DegValAlaGluIleGlnLeuMetHisGlnHarAlaLysTrp (SEQ. ID. NO. 24);
- 10 (b) peptides containing amino acids 1-9, 1-10, 1-11, 1-12 or 1-13;
- (c) pharmaceutically acceptable salts of (a) or (b); or
- (d) N - or C - derivatives of (a), (b) or (c).
6. The peptide of claim 1, wherein said peptide is selected from:
- (a) DegValDegGluIleGlnLeuMetHisGlnHarAlaLysTrp (SEQ. ID. NO. 27);
- 15 (b) peptides containing amino acids 1-9, 1-10, 1-11, 1-12 or 1-13;
- (c) pharmaceutically acceptable salts of (a) or (b); or
- (d) N - or C - derivatives of (a), (b) or (c).
7. The peptide of claim 1, wherein said peptide is selected from:
- (a) DegValAc₃cGluIleGlnLeuMetHisGlnHarAlaLysTrp (SEQ. ID. NO. 40);
- 20 (b) peptides containing amino acids 1-9, 1-10, 1-11, 1-12 or 1-13;
- (c) pharmaceutically acceptable salts of (a) or (b); or
- 25 (d) N - or C - derivatives of (a), (b) or (c).

-38-

8. The peptide of claim 1, wherein said peptide is selected from:
- (a) DegValAc₃cGluIleGlnLeuMetHisGlnHarAlaLysTrp (SEQ. ID. NO. 41);
 - (b) peptides containing amino acids 1-9, 1-10, 1-11, 1-12 or 1-13;
 - (c) pharmaceutically acceptable salts of (a) or (b); or
 - (d) N - or C - derivatives of (a), (b) or (c).
9. The peptide of claim 1, wherein said peptide is selected from:
- (a) DegValAibGluIleGlnLeuMetHisGlnHarAlaLysTrp (SEQ. ID. NO. 42);
 - (b) peptides containing amino acids 1-9, 1-10, 1-11, 1-12 or 1-13;
 - (c) pharmaceutically acceptable salts of (a) or (b); or
 - (d) N - or C - derivatives of (a), (b) or (c).
10. The peptide of claim 1, wherein said peptide is selected from:
- (a) Ac₃cValAlaGluIleGlnLeuMetHisGlnHarAlaLysTrp (SEQ. ID. NO. 25);
 - (b) peptides containing amino acids 1-9, 1-10, 1-11, 1-12 or 1-13;
 - (c) pharmaceutically acceptable salts of (a) or (b); or
 - (d) N - or C - derivatives of (a), (b) or (c).
11. The peptide of claim 1, wherein said peptide is selected from:
- (a) Ac₃cValDegGluIleGlnLeuMetHisGlnHarAlaLysTrp (SEQ. ID. NO. 43);
 - (b) peptides containing amino acids 1-9, 1-10, 1-11, 1-12 or 1-13;
 - (c) pharmaceutically acceptable salts of (a) or (b); or
 - (d) N - or C - derivatives of (a), (b) or (c).
12. The peptide of claim 1, wherein said peptide is selected from:
- (a) Ac₃cValAc₃cGluIleGlnLeuMetHisGlnHarAlaLysTrp (SEQ. ID. NO. 28);

-39-

- (b) peptides containing amino acids 1-9, 1-10, 1-11, 1-12 or 1-13;
- (c) pharmaceutically acceptable salts (a) or (b); or
- (d) N - or C - derivatives of (a), (b) or (c).

13. The peptide of claim 1, wherein said peptide is selected from:

- 5 (a) $\text{Ac}_3\text{cValAc}_3\text{cGluIleGlnLeuMetHisGlnHarAlaLysTrp}$ (SEQ. ID. NO. 44);
- (b) peptides containing amino acids 1-9, 1-10, 1-11, 1-12 1-13;
 - (c) pharmaceutically acceptable salts (a) or (b); or
 - (d) N - or C - derivatives of (a), (b) or (c).

14. The peptide of claim 1, wherein said peptide is selected from:

- 10 (a) $\text{Ac}_3\text{cValAibGluIleGlnLeuMetHisGlnHarAlaLysTrp}$ (SEQ. ID. NO. 45);
- (b) peptides containing amino acids 1-9, 1-10, 1-11, 1-12 or 1-13;
 - (c) pharmaceutically acceptable salts of (a) or (b); or
 - 15 (d) N - or C - derivatives of (a), (b) or (c).

15. The peptide of claim 1, wherein said peptide is selected from:

- (a) $\text{Ac}_3\text{cValAlaGluIleGlnLeuMetHisGlnHarAlaLysTrp}$ (SEQ. ID. NO. 4);
- (b) peptides containing amino acids 1-9, 1-10, 1-11, 1-12 or 1-13;
 - 20 (c) pharmaceutically acceptable salts of (a) or (b); or
 - (d) N - or C - derivatives of (a), (b) or (c).

16. The peptide of claim 1, wherein said peptide is selected from:

- (a) $\text{Ac}_3\text{cValDegGluIleGlnLeuMetHisGlnHarAlaLysTrp}$ (SEQ. ID. NO. 46);
- 25 (b) peptides containing amino acids 1-9,1-10, 1-11, 1-12 or 1-13;
 - (c) pharmaceutically acceptable salts of (a) or (b); or
 - (d) N - or C - derivatives of (a), (b) or (c).

-40-

17. The peptide of claim 1, wherein said peptide is selected from:
- (a) Ac₃cValAc₃cGluIleGlnLeuMetHisGlnHarAlaLysTrp (SEQ. ID. NO. 47);
- 5 (b) peptides containing amino acids 1-9, 1-10, 1-11, 1-12 or 1-13;
- (c) pharmaceutically acceptable salts of (a) or (b); or
- (d) N - or C - derivatives of (a), (b) or (c).
18. The peptide of claim 1, wherein said peptide is selected from:
- (a) Ac₃cValAc₃cGluIleGlnLeuMetHisGlnHarAlaLysTrp (SEQ. ID. NO. 29);
- 10 (b) peptides containing amino acids 1-9, 1-10, 1-11, 1-12 or 1-13;
- (c) pharmaceutically acceptable salts of (a) or (b); or
- (d) N - or C - derivatives of (a), (b) or (c).
19. The peptide of claim 1, wherein said peptide is selected from:
- (a) Ac₃cValAibGluIleGlnLeuMetHisGlnHarAlaLysTrp (SEQ. ID. NO. 15);
- 15 (b) peptides containing amino acids 1-9, 1-10, 1-11, 1-12, or 1-13;
- (c) pharmaceutically acceptable salts of (a) or (b); or
- (d) N - or C - derivatives of (a), (b) or (c).
20. The peptide of claim 1, wherein said peptide is selected from:
- (a) AibValDegGluIleGlnLeuMetHisGlnHarAlaLysTrp (SEQ. ID. NO. 48);
- 20 (b) peptides containing amino acids 1-9, 1-10, 1-11, 1-12 or 1-13;
- (c) pharmaceutically acceptable salts of (a) or (b); or
- (d) N - or C - derivatives of (a), (b) or (c).
21. The peptide of claim 1, wherein said peptide is selected from:
- (a) AibValAc₃cGluIleGlnLeuMetHisGlnHarAlaLysTrp (SEQ. ID. NO. 49);
- 25

-41-

- (b) peptides containing amino acids 1-9, 1-10, 1-11, 1-12 or 1-13;
 - (c) pharmaceutically acceptable salts of (a) or (b); or
 - (d) N - or C - derivatives of (a), (b) or (c).
22. The peptide of claim 1, wherein said peptide is selected from:
- 5 (a) AibValAc₅cGluIleGlnLeuMetHisGlnHarAlaLysTrp (SEQ. ID. NO. 50);
- (b) peptides containing amino acids 1-9, 1-10, 1-11, 1-12 or 1-13;
 - (c) pharmaceutically acceptable salts of (a) or (b); or
 - (d) N - or C - derivatives of (a), (b) or (c).
- 10 23. The peptide of claim 1, wherein said peptide is selected from:
- (a) Ac₅cValSerGluIleGlnLeuMetHisGlnHarAlaLysTrp (SEQ. ID. NO. 51);
- (b) peptides containing amino acids 1-9, 1-10, 1-11, 1-12 or 1-13;
 - (c) pharmaceutically acceptable salts of (a) or (b); or
 - 15 (d) N - or C - derivatives of (a), (b) or (c).
24. The peptide of claim 1, wherein said peptide is selected from:
- (a) Ac₅cValSerGluIleGlnLeuMetHisAsnLeuGlyLysHis (SEQ. ID. NO. 52);
- (b) peptides containing amino acids 1-9, 1-10, 1-11, 1-12, or 1-13;
 - 20 (c) pharmaceutically acceptable salts of (a) or (b); or
 - (d) N - or C - derivatives of (a), (b) or (c).
25. The peptide of claim 1, wherein said peptide is selected from:
- (a) Ac₅cValAlaGluIleGlnLeuMetHis (part of SEQ. ID. NO. 4);
 - (b) pharmaceutically acceptable salts thereof; or
 - 25 (c) N - or C - derivatives of (a) or (b).
26. A biologically active peptide consisting essentially of the formula selected

-42-

from:

- (a) X_{01} Val X_{02} GluIle X_{03} LeuMetHis X_{04} X_{05} X_{06} Lys X_{07} (SEQ. ID. NO. 5);
- (b) peptides containing amino acids 1-9, 1-10, 1-11, 1-12, or 1-13;
- (c) pharmaceutically acceptable salts of (a) or (b); or
- (d) N - or C - derivatives of (a), (b) or (c);

wherein:

X_{01} is α -helix-stabilizing residue, Gly, Ser or Ala;

X_{02} is α -helix-stabilizing residue, Ala or Ser;

X_{03} is Ala, Gln or Asn;

X_{04} is Ala, Gln, Asn, Har or Arg;

X_{05} is an α -helix stabilizing residue, Ala or Gly;

X_{06} is an α -helix stabilizing residue or Lys;

X_{07} is α -helix stabilizing residue, Trp, or His;

wherein at least one of X_{01} , X_{02} , X_{05} , X_{06} or X_{07} is an α -helix stabilizing residue,

and wherein at least one of said α -helix stabilizing residues is Aib, Ac₃c, Ac₄c, Ac₅c, Ac₆c, or Deg.

27. The peptide of claim 26, wherein said peptide is selected from:

- (a) Ac₄cValAibGluIleGlnLeuMetHisGlnHarAlaLysTrp (SEQ. ID. NO. 7);
- (b) peptides containing amino acids 1-9, 1-10, 1-11, 1-12 or 1-13;
- (c) pharmaceutically acceptable salts of (a) or (b); or
- (d) N - or C - derivatives (a), (b) or (c).

28. The peptide of claim 26, wherein said peptide is selected from:

- (a) Ac₆cValAibGluIleGlnLeuMetHisGlnHarAlaLysTrp (SEQ. ID. NO. 8);
- (b) peptides containing amino acids 1-9, 1-10, 1-11, 1-12 or 1-13;
- (c) pharmaceutically acceptable salts of (a) or (b); or
- (d) N - or C - derivatives of (a), (b) or (c).

-43-

29. The peptide of claim 26, wherein said peptide is selected from:
- (a) $\text{Ac}_5\text{cVal Ac}_4\text{cGluIleGlnLeuMetHisGlnHarAlaLysTrp}$ (SEQ. ID. NO. 9);
 - (b) peptides containing amino acids 1-9, 1-10, 1-11, 1-12 or 1-13;
 - (c) pharmaceutically acceptable salts of (a) or (b); or
 - (d) N - or C - derivatives (a), (b) or (c).
30. The peptide of claim 26, wherein said peptide is selected from:
- (a) $\text{Ac}_5\text{cValAc}_6\text{cGluIleGlnLeuMetHisGlnHarAlaLysTrp}$ (SEQ. ID. NO. 10);
 - (b) peptides containing amino acids 1-9, 1-10, 1-11, 1-12 or 1-13;
 - (c) pharmaceutically acceptable salts of (a) or (b); or
 - (d) N - or C - derivatives of (a), (b) or (c).
31. The peptide of claim 26, wherein said peptide is selected from:
- (a) $\text{Ac}_4\text{cValAc}_4\text{cGluIleGlnLeuMetHisGlnHarAlaLysTrp}$ (SEQ. ID. NO. 11);
 - (b) peptides containing amino acids 1-9, 1-10, 1-11, 1-12 or 1-13;
 - (c) pharmaceutically acceptable salts of (a) or (b); or
 - (d) N - or C - derivatives of (a), (b) or (c).
32. The peptide of claim 26, wherein said peptide is selected from:
- (a) $\text{Ac}_6\text{cValAc}_6\text{cGluIleGlnLeuMetHisGlnHarAlaLysTrp}$ (SEQ. ID. NO. 12);
 - (b) peptides containing amino acids 1-9, 1-10, 1-11, 1-12 or 1-13;
 - (c) pharmaceutically acceptable salts of (a) or (b); or
 - (d) N - or C - derivatives of (a), (b) or (c).
33. The peptide of claim 1 or 26, wherein said peptide is labeled with a label selected from the group consisting of a fluorescent label, a chemiluminescent label, a bioluminescent label and a radioactive label.

-44-

34. The peptide of claim 1 or 26, wherein said peptide is labeled with ^{125}I .
35. The peptide of claim 1 or 26, wherein said peptide is labeled with $^{99\text{m}}\text{Tc}$.
36. A pharmaceutical composition comprising the biologically active peptide of claim 1 or 26, and a pharmaceutically acceptable carrier.
- 5 37. A method for treating mammalian conditions characterized by decreases in bone mass, said method comprising administering to a subject in need thereof an effective bone-mass increasing amount of a biologically active peptide of claim 1 or 26.
- 10 38. A method for treating mammalian conditions characterized by decreases in bone mass, said method comprising administering to a subject in need thereof an effective bone mass-increasing amount of a composition comprising a biologically active peptide of claim 1 or 26 and a pharmaceutically acceptable carrier.
- 15 39. A method for determining rates of bone reformation, bone resorption and/or bone remodeling comprising administering to a patient an effective amount of a peptide of claim 1 or 26 and determining the uptake of said peptide into the bone of said patient.
40. The method of claim 37, wherein said condition to be treated is hyperparathyroidism.
- 20 41. The method of claim 37, wherein said condition to be treated is hypercalcemia.
42. The method of claim 37, wherein said effective amount of said peptide for increasing bone mass is from about $0.01\mu\text{g/kg/day}$ to about $1.0\mu\text{g/kg/day}$.

-45-

43. The method of claim 37, wherein the method of administration is parenteral.

44. The method of claim 37, wherein the method of administration is subcutaneous.

5 45. The method of claim 37, wherein the method of administration is nasal insufflation.

46. The method of claim 37, wherein the method of administration is oral.

47. The method of making the peptide of claim 1 or 26, wherein said peptide is synthesized by solid phase synthesis.

10 48. The method of making the peptide of claim 1 or 26, wherein said peptide is synthesized by liquid phase synthesis.

49. The method of making the peptide of claim 1 or 26, wherein said peptide is protected by FMOC.

1/9

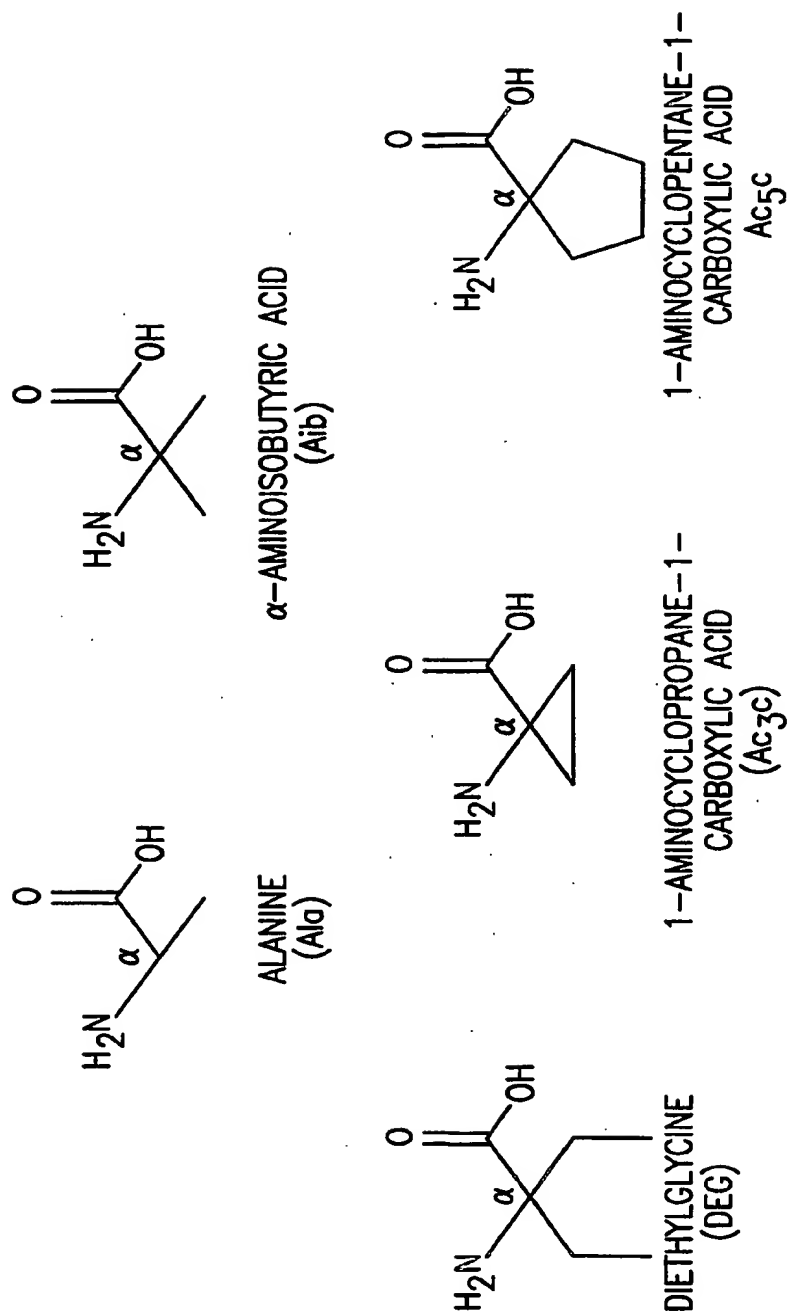


FIG.1

2/9

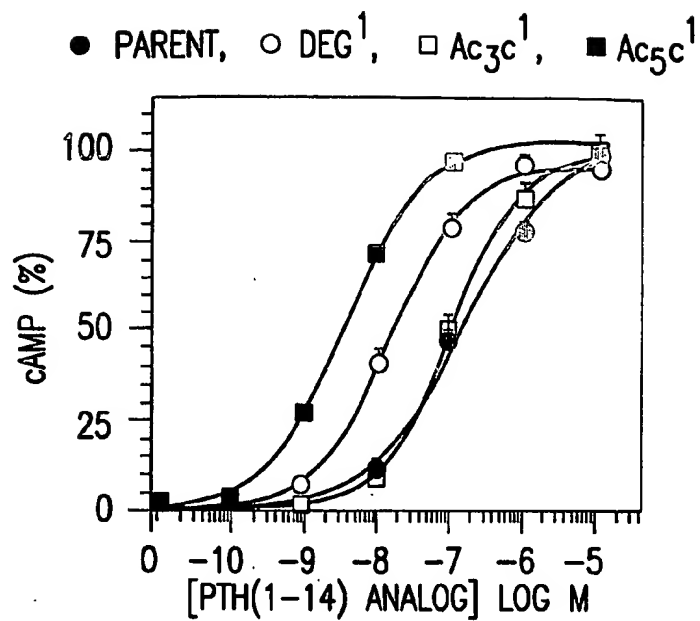


FIG.2A

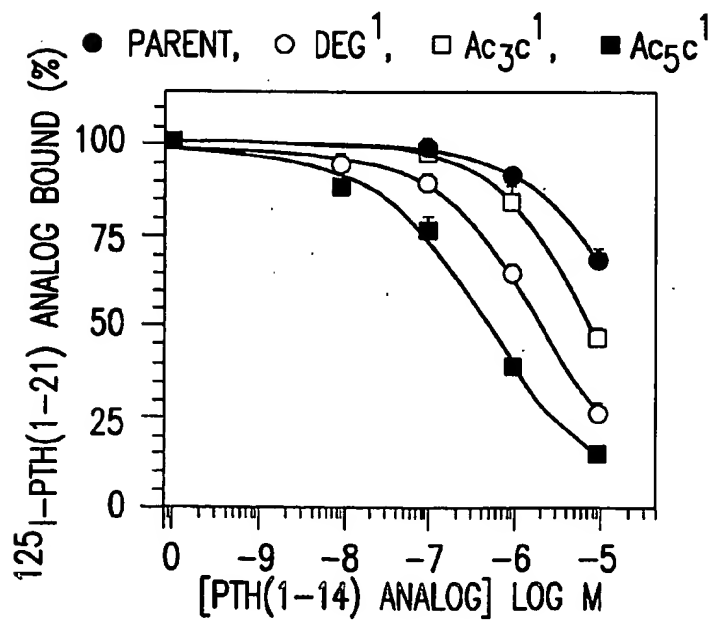


FIG.2B

3/9

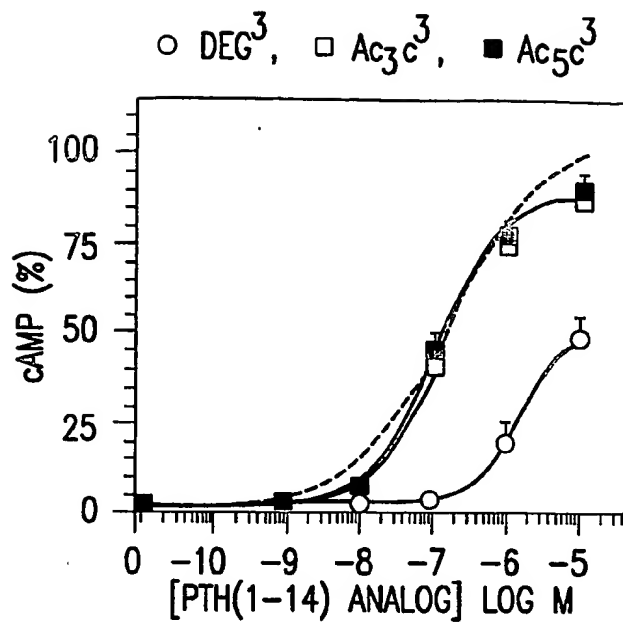


FIG.2C

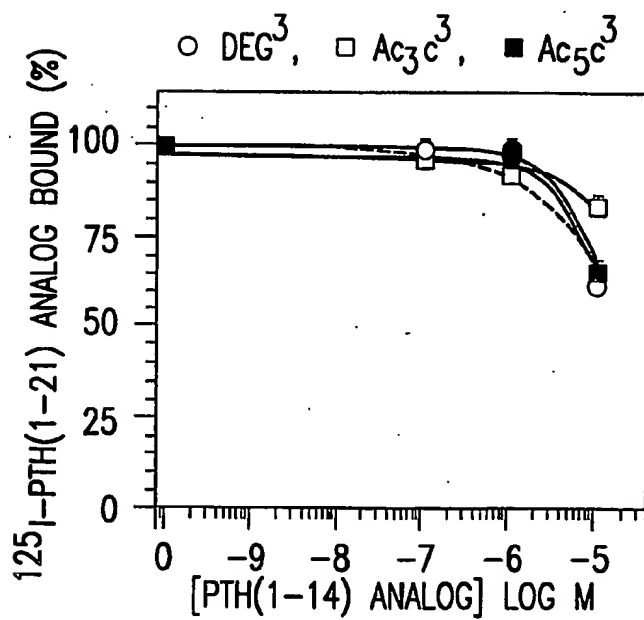


FIG.2D

4/9

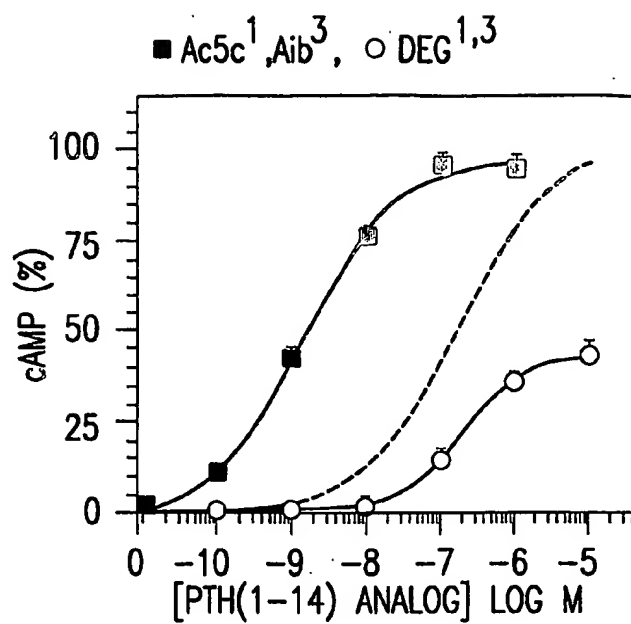


FIG.3A

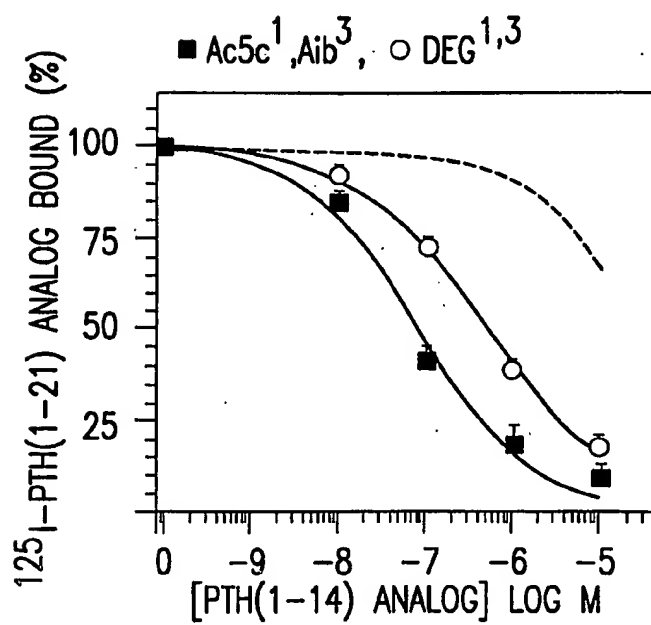


FIG.3B

5/9

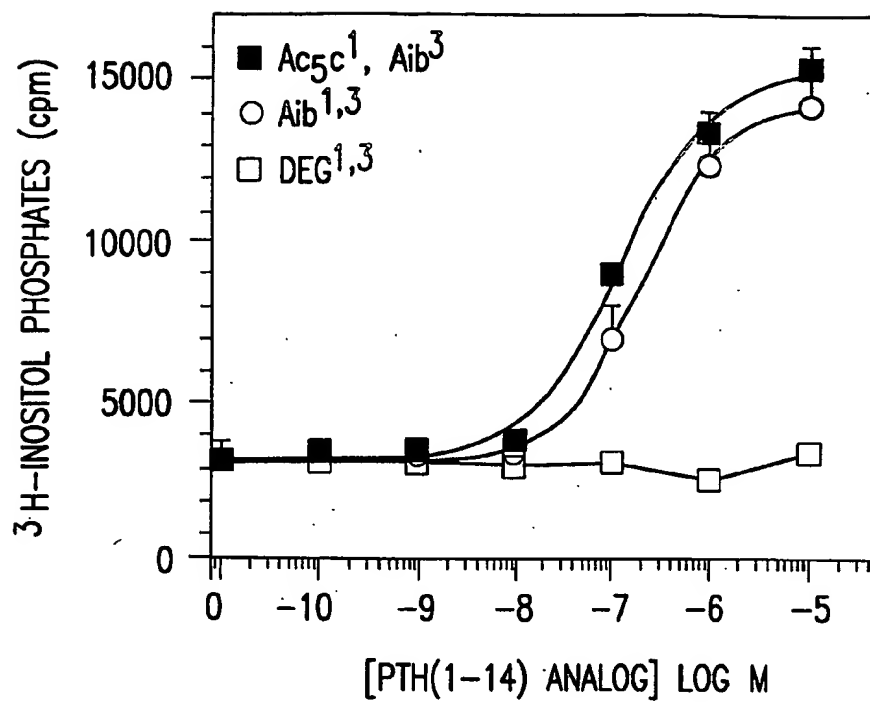


FIG.4

6/9

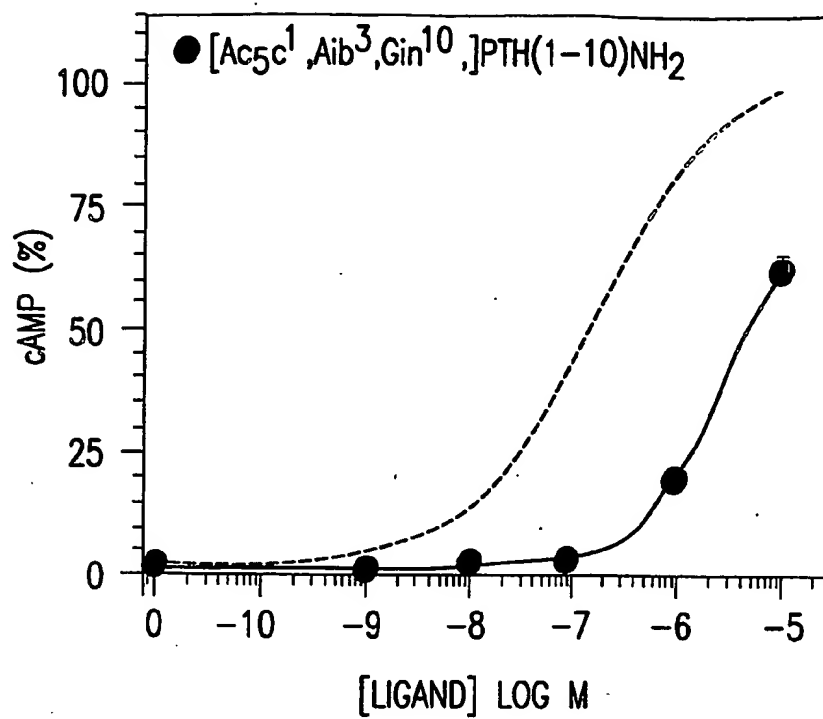


FIG.5A

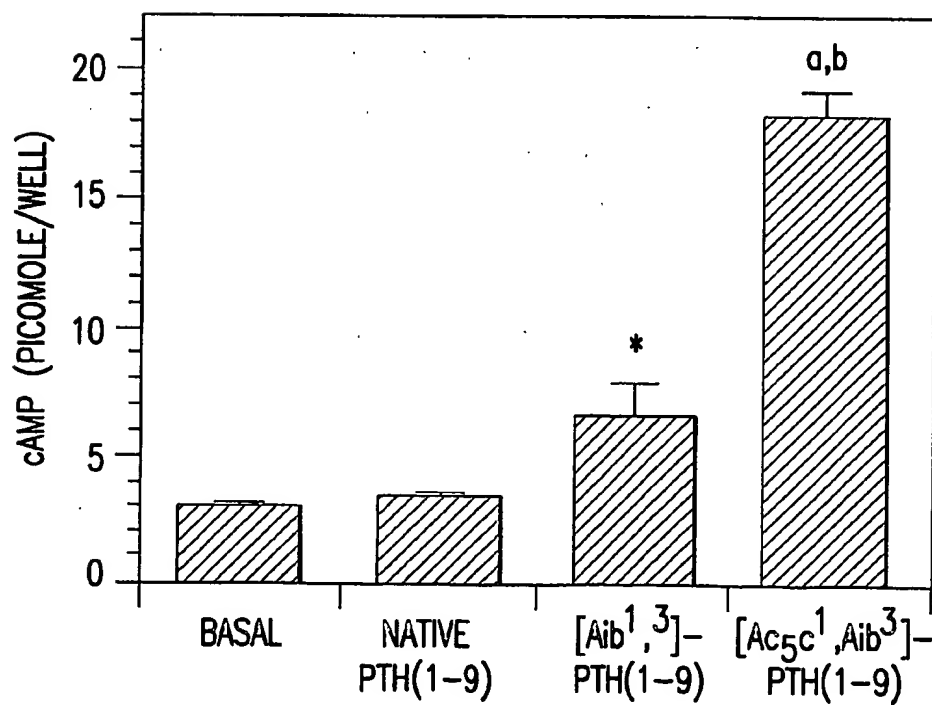


FIG.5B

7/9

- [Aib^{1,3},Gln¹⁰,Har¹¹,Ala¹²,Trp¹⁴]PTH(1-14)NH₂
○ Ac₅c¹,Aib³,Gln¹⁰,Har¹¹,Ala¹²,Trp¹⁴]PTH(1-14)NH₂
□ Deg^{1,3},Gln¹⁰,Har¹¹,Ala¹²,Trp¹⁴]PTH(1-14)NH₂
△ [Ac₅c¹,Aib³,Gln¹⁰,]PTH(1-10)NH₂

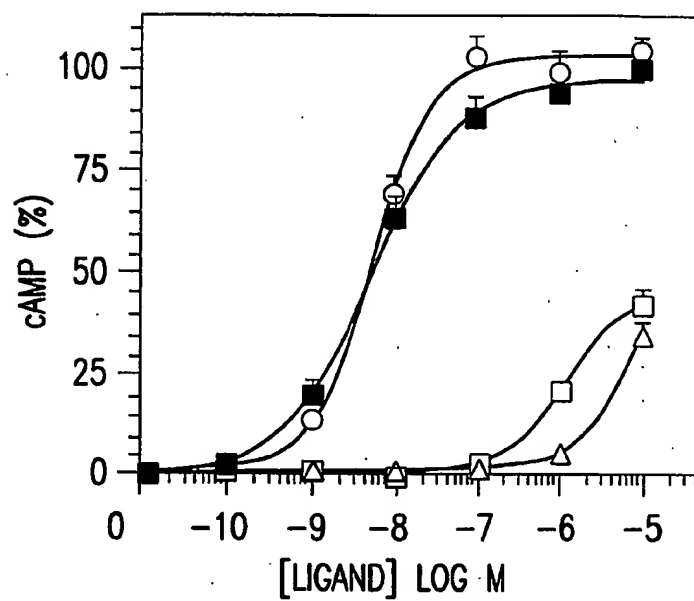


FIG.6

8/9

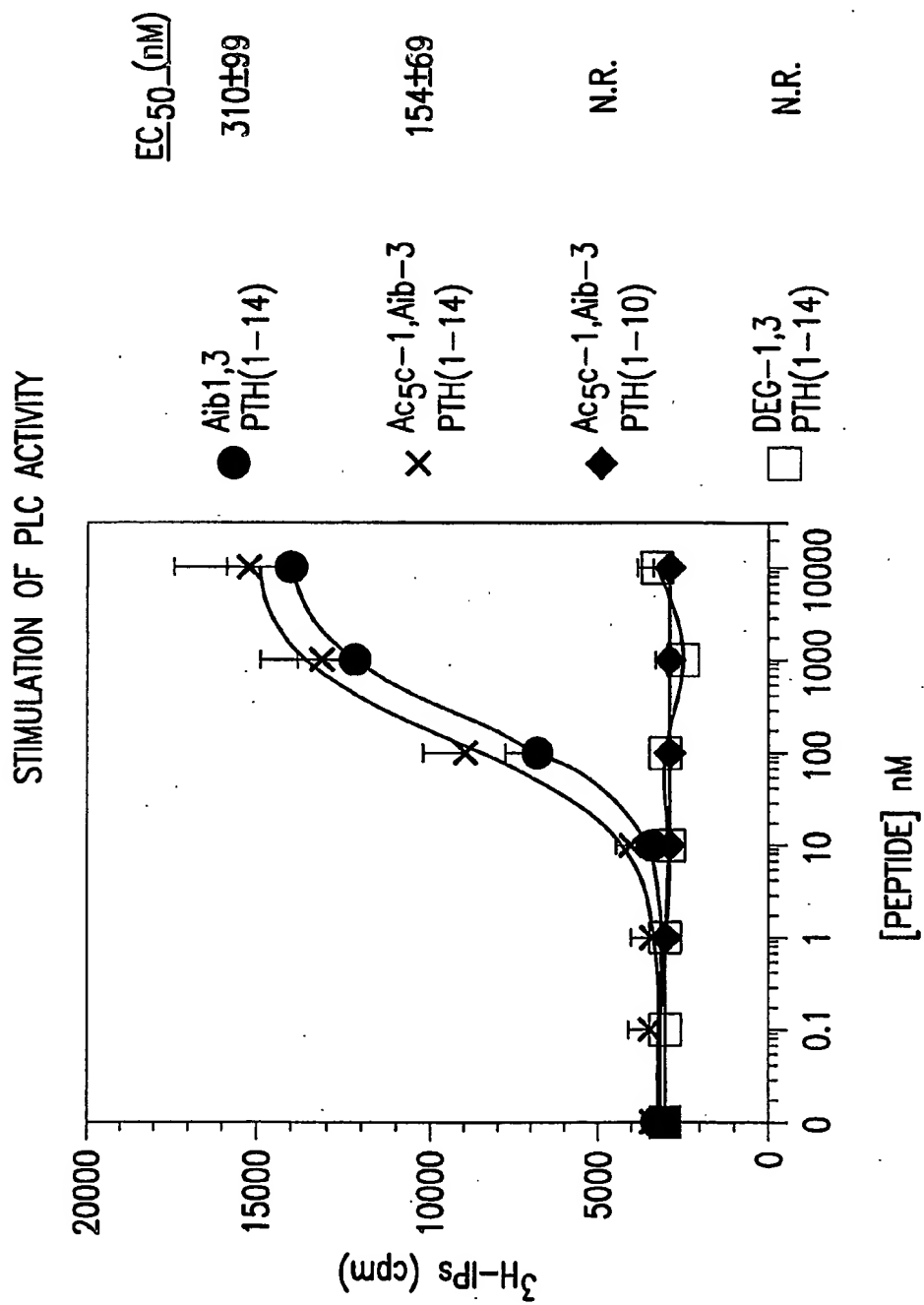


FIG.7

9/9

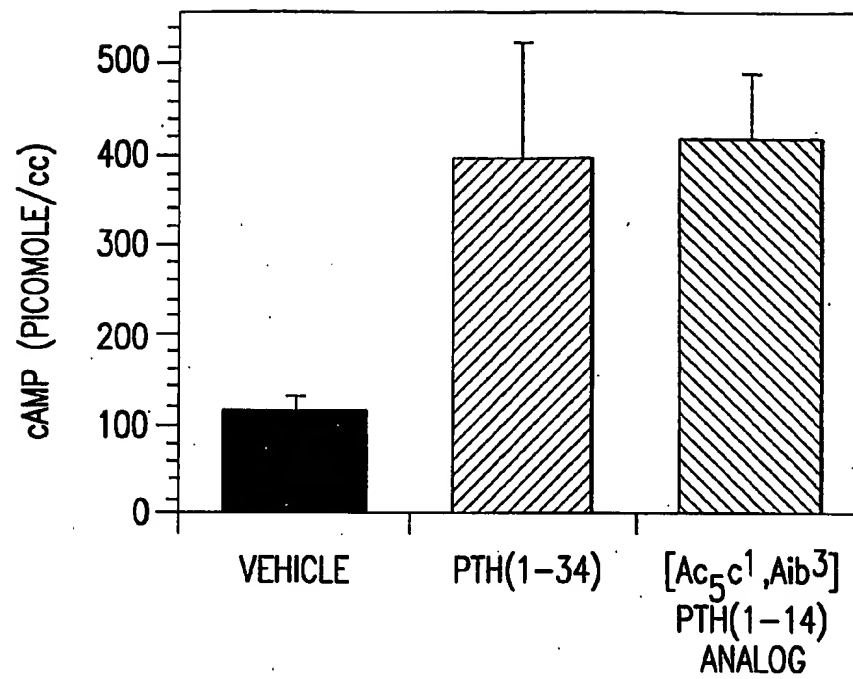


FIG.8

-1-

SEQUENCE LISTING

<110> Gardella, PhD, Thomas J.
Potts, John T.
Kronenberg, H.M.
Shimizu, N.
Carter, P.

<120> Conformationally Constrained Parathyroid Hormones With
Alpha-Helix Stabilizers

<130> 0609.515PC00

<160> 52

<170> PatentIn version 3.2

<210> 1
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> Mutated hPTH

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> Xaa can represent an alpha-helix-stabilizing residue, Gly, Ser or
Ala

<220>
<221> MISC_FEATURE
<222> (1)..(14)
<223> At least one Xaa is an alpha-helix-stabilizing residue

<220>
<221> MISC_FEATURE
<222> (1)..(14)
<223> At least one Xaa is 1-aminocyclopropane-1-carboxylic acid,
1-amino-cyclobutane-carboxylic acid,
1-aminocyclopentane-1-carboxylic acid,
1-amino-cyclohexane-carboxylic acid or alpha, alpha-diethylglycine

<220>
<221> MISC_FEATURE
<222> (1)..(14)
<223> At least one Xaa is 1-aminocyclopropane-1-carboxylic acid,
1-amino-cyclobutane-carboxylic acid,
1-aminocyclopentane-1-carboxylic acid, 1-amino-cyclohexane-
carboxylic acid or alpha, alpha-diethylglycine

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa can represent an alpha-helix-stabilizing residue, Ala or Ser

<220>
<221> MISC_FEATURE
<222> (10)..(10)

-2-

<223> Xaa can represent Ala, Gln or Asn

<220>

<221> MISC_FEATURE

<222> (11)..(11)

<223> Xaa can represent Arg, Har or Leu

<220>

<221> MISC_FEATURE

<222> (12)..(12)

<223> Xaa can represent alpha-helix-stabilizing residue, Ala or Gly

<220>

<221> MISC_FEATURE

<222> (13)..(13)

<223> Xaa can represent alpha-helix-stabilizing residue or Lys

<220>

<221> MISC_FEATURE

<222> (14)..(14)

<223> Xaa can represent alpha-helix-stabilizing residue, Trp or His

<220>

<221> MOD_RES

<222> (14)..(14)

<223> AMIDATION

<400> 1

Xaa	Val	Xaa	Glu	Ile	Gln	Leu	Met	His	Xaa	Xaa	Xaa	Xaa	Xaa
1				5					10				

<210> 2

<211> 14

<212> PRT

<213> Artificial Sequence

<220>

<223> Mutated hPTH

<220>

<221> MISC_FEATURE

<222> (1)..(1)

<223> Xaa can represent an alpha-stabilizing residue, Gly, Ser or Ala

<220>

<221> MISC_FEATURE

<222> (1)..(3)

<223> At least one Xaa is an alpha-helix-stabilizing residue

<220>

<221> MISC_FEATURE

<222> (1)..(3)

<223> At least one Xaa is 1-aminocyclopropane-1-carboxylic acid,
1-amino-cyclobutane-carboxylic acid,
1-aminocyclopentane-1-carboxylic acid,
1-amino-cyclohexane-carboxylic acid or alpha, alpha-diethylglycine

<220>

-3-

<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa can represent an alpha-stabilizing residue, Ala or Ser

<220>
<221> MISC_FEATURE
<222> (11)..(11)
<223> Xaa represents homoarginine

<220>
<221> MOD_RES
<222> (14)..(14)
<223> AMIDATION

<400> 2

Xaa Val Xaa Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp
1 5 10

<210> 3
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> [Aib-1,3, M]PTH(1-14)

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> Xaa represents alpha-aminoisobutyric acid

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa represents alpha-aminoisobutyric acid

<220>
<221> MISC_FEATURE
<222> (11)..(11)
<223> Xaa represents homoarginine

<220>
<221> MOD_RES
<222> (14)..(14)
<223> AMIDATION

<400> 3

Xaa Val Xaa Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp
1 5 10

<210> 4
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> [Ac5c-1, M]PTH(1-14)

-4-

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> Xaa represents 1-aminocyclopentane-1-carboxylic acid

<220>
<221> MISC_FEATURE
<222> (11)..(11)
<223> Xaa represents homoarginine

<220>
<221> MOD_RES
<222> (14)..(14)
<223> AMIDATION

<400> 4

Xaa Val Ala Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp
1 5 10

<210> 5
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> Mutated hPTH

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> Xaa can represent an alpha-helix-stabilizing residue, Gly, Ser or Ala

<220>
<221> MISC_FEATURE
<222> (1)..(14)
<223> At least one Xaa is an alpha-helix-stabilizing residue

<220>
<221> MISC_FEATURE
<222> (1)..(14)
<223> At least one Xaa is 1-aminocyclopropane-1-carboxylic acid, 1-amino-cyclobutane-carboxylic acid, 1-aminocyclopentane-1-carboxylic acid, 1-amino-cyclohexane-carboxylic acid or alpha, alpha-diethylglycine

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa can represent alpha-helix-stabilizing residue, Ala or Ser

<220>
<221> MISC_FEATURE
<222> (6)..(6)
<223> Xaa can represent Ala, Gln or Asn

<220>
<221> MISC_FEATURE

-5-

<222> (10)..(10)
 <223> Xaa can represent Ala, Gln or Asn

 <220>
 <221> MISC_FEATURE
 <222> (11)..(11)
 <223> Xaa can represent an alpha-helix-stabilizing residue, Har or Arg

 <220>
 <221> MISC_FEATURE
 <222> (12)..(12)
 <223> Xaa can represent an alpha-helix-stabilizing residue or Lys

 <220>
 <221> MISC_FEATURE
 <222> (14)..(14)
 <223> Xaa can represent an alpha-helix-stabilizing residue, Trp or His

 <220>
 <221> MOD_RES
 <222> (14)..(14)
 <223> AMIDATION

 <400> 5

Xaa Val Xaa Glu Ile Xaa Leu Met His Xaa Xaa Xaa Lys Xaa
 1 5 10

<210> 6
 <211> 14
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Mutated hPTH

<220>
 <221> MISC_FEATURE
 <222> (1)..(1)
 <223> Xaa can represent an alpha-helix-stabilizing residue, Gly, Ser or Ala

<220>
 <221> MISC_FEATURE
 <222> (1)..(14)
 <223> At least one Xaa is 1-aminocyclopropane-1-carboxylic acid, 1-amino-cyclobutane-carboxylic acid, 1-aminocyclopentane-1-carboxylic acid, 1-amino-cyclohexane-carboxylic acid or alpha, alpha-diethylglycine

<220>
 <221> MISC_FEATURE
 <222> (3)..(3)
 <223> Xaa can represent an alpha-helix-stabilizing residue, Ala or Ser

<220>
 <221> MISC_FEATURE
 <222> (11)..(11)
 <223> Xaa represents homoarginine

-6-

<220>
<221> MOD_RES
<222> (14)..(14)
<223> AMIDATION

<400> 6

Xaa Val Xaa Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp
1 5 10

<210> 7
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> [Ac4c-1, Aib-3, M]PTH(1-14)

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> Xaa represents 1-amino-cyclobutane-carboxylic acid

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa represents alpha-aminoisobutyric acid

<220>
<221> MISC_FEATURE
<222> (11)..(11)
<223> Xaa represents homoarginine

<220>
<221> MOD_RES
<222> (14)..(14)
<223> AMIDATION

<400> 7

Xaa Val Xaa Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp
1 5 10

<210> 8
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> [Ac6c-1, Aib-3, M]PTH(1-14)

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> Xaa represents 1-amino-cyclohexane-carboxylic acid

<220>
<221> MISC_FEATURE

-7-

<222> (3)..(3)
<223> Xaa represents alpha-aminoisobutyric acid

<220>
<221> MISC_FEATURE
<222> (11)..(11)
<223> Xaa represents homoarginine

<220>
<221> MOD_RES
<222> (14)..(14)
<223> AMIDATION

<400> 8

Xaa Val Xaa Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp
1 5 10

<210> 9
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> [Ac5c-1, Ac4c-3, M]PTH(1-14)

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> Xaa represents 1-aminocyclopentane-1-carboxylic acid

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa represents 1-amino-cyclobutane-carboxylic acid

<220>
<221> MISC_FEATURE
<222> (11)..(11)
<223> Xaa represents homoarginine

<220>
<221> MOD_RES
<222> (14)..(14)
<223> AMIDATION

<400> 9

Xaa Val Xaa Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp
1 5 10

<210> 10
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> [Ac5c-1, Ac6c-3, M]PTH(1-14)

-8-

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> Xaa represents 1-aminocyclopentane-1-carboxylic acid

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa represents 1-amino-cyclohexane-carboxylic acid

<220>
<221> MISC_FEATURE
<222> (11)..(11)
<223> Xaa represents homoarginine

<220>
<221> MOD_RES
<222> (14)..(14)
<223> AMIDATION

<400> 10

Xaa Val Xaa Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp
1 5 10

<210> 11
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> [Ac4c-1,3, M]PTH(1-14)

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> Xaa represents 1-amino-cyclobutane-carboxylic acid

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa represents 1-amino-cyclobutane-carboxylic acid

<220>
<221> MISC_FEATURE
<222> (11)..(11)
<223> Xaa represents homoarginine

<220>
<221> MOD_RES
<222> (14)..(14)
<223> AMIDATION

<400> 11

Xaa Val Xaa Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp
1 5 10

<210> 12

-9-

<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> [Ac6c-1,3, M] PTH(1-14)

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> Xaa represents 1-amino-cyclohexane-carboxylic acid

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa represents 1-amino-cyclohexane-carboxylic acid

<220>
<221> MISC_FEATURE
<222> (11)..(11)
<223> Xaa represents homoarginine

<220>
<221> MOD_RES
<222> (14)..(14)
<223> AMIDATION

<400> 12

Xaa Val Xaa Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp
1 5 10

<210> 13
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> [M] PTH(1-14)

<220>
<221> MISC_FEATURE
<222> (11)..(11)
<223> Xaa represents homoarginine

<220>
<221> MOD_RES
<222> (14)..(14)
<223> AMIDATION

<400> 13

Ala Val Ala Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp
1 5 10

<210> 14
<211> 34
<212> PRT

-10-

<213> Artificial Sequence

<220>

<223> [Nle-8,21, Tyr-34]rPTH(1-34)amide

<220>

<221> MISC_FEATURE

<222> (8)..(8)

<223> Xaa represents Norleucine

<220>

<221> MISC_FEATURE

<222> (21)..(21)

<223> Xaa represents Norleucine

<220>

<221> MOD_RES

<222> (34)..(34)

<223> AMIDATION

<400> 14

Ala	Val	Ser	Glu	Ile	Gln	Leu	Xaa	His	Asn	Leu	Gly	Lys	His	Leu	Ala
1				5					10					15	

Ser	Val	Glu	Arg	Xaa	Gln	Trp	Leu	Arg	Lys	Lys	Leu	Gln	Asp	Val	His
			20					25					30		

Asn Tyr

<210> 15

<211> 14

<212> PRT

<213> Artificial Sequence

<220>

<223> [Ac5c-1, Aib-3, M]PTH(1-14)amide

<220>

<221> MISC_FEATURE

<222> (1)..(1)

<223> Xaa represents 1-aminocyclopentane-1-carboxylic acid

<220>

<221> MISC_FEATURE

<222> (3)..(3)

<223> Xaa represents alpha-aminoisobutyric acid

<220>

<221> MISC_FEATURE

<222> (11)..(11)

<223> Xaa represents homoarginine

<220>

<221> MOD_RES

<222> (14)..(14)

-11-

<223> AMIDATION

<400> 15

Xaa Val Xaa Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp
1 5 10

<210> 16

<211> 11

<212> PRT

<213> Artificial Sequence

<220>

<223> [Ala-1,3, Gln-10, Har-11]PTH(1-11)amide

<220>

<221> MISC_FEATURE

<222> (11)..(11)

<223> Xaa represents homoarginine

<220>

<221> MOD_RES

<222> (11)..(11)

<223> AMIDATION

<400> 16

Ala Val Ala Glu Ile Gln Leu Met His Gln Xaa
1 5 10

<210> 17

<211> 14

<212> PRT

<213> Rattus sp.

<220>

<221> MOD_RES

<222> (14)..(14)

<223> AMIDATION

<400> 17

Ala Val Ser Glu Ile Gln Leu Met His Asn Leu Gly Lys His
1 5 10

<210> 18

<211> 84

<212> PRT

<213> Homo sapiens

<220>

<221> MOD_RES

<222> (84)..(84)

<223> AMIDATION

<400> 18

-12-

Ser Val Ser Glu Ile Gln Leu Met His Asn Leu Gly Lys His Leu Asn
1 5 10 15

Ser Met Glu Arg Val Glu Trp Leu Arg Lys Lys Leu Gln Asp Val His
20 25 30

Asn Phe Val Ala Leu Gly Ala Pro Leu Ala Pro Arg Asp Ala Gly Ser
35 40 45

Gln Arg Pro Arg Lys Lys Glu Asp Asn Val Leu Val Glu Ser His Glu
50 55 60

Lys Ser Leu Gly Glu Ala Asp Lys Ala Asp Val Asn Val Leu Thr Lys
65 70 75 80

Ala Lys Ser Gln

<210> 19
<211> 34
<212> PRT
<213> Homo sapiens

<220>
<221> MOD_RES
<222> (34)..(34)
<223> AMIDATION

<400> 19

Ser Val Ser Glu Ile Gln Leu Met His Asn Leu Gly Lys His Leu Asn
1 5 10 15

Ser Met Glu Arg Val Glu Trp Leu Arg Lys Lys Leu Gln Asp Val His
20 25 30

Asn Phe

<210> 20
<211> 21
<212> PRT
<213> Artificial Sequence

<220>
<223> [M]PTH(1-21)

<220>
<221> MISC_FEATURE
<222> (11)..(11)
<223> Xaa represents homoarginine

-13-

<220>
<221> MOD_RES
<222> (21)..(21)
<223> AMIDATION

<400> 20

Ala Val Ala Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp Leu Asn
1 5 10 15

Ser Met Glu Arg Val
20

<210> 21
<211> 21
<212> PRT
<213> Artificial Sequence

<220>
<223> [Aib-1,3, M] PTH(1-21)

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> Xaa represents alpha-aminoisobutyric acid

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa represents alpha-aminoisobutyric acid

<220>
<221> MISC_FEATURE
<222> (11)..(11)
<223> Xaa represents homoarginine

<220>
<221> MOD_RES
<222> (21)..(21)
<223> AMIDATION

<400> 21

Xaa Val Xaa Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp Leu Asn
1 5 10 15

Ser Met Glu Arg Val
20

<210> 22
<211> 10
<212> PRT
<213> Artificial Sequence

<220>
<223> [Ac5c-1, Aib-3, Gln-10] PTH(1-10)

-14-

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> Xaa represents 1-aminocyclopentane-1-carboxylic acid

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa represents alpha-aminoisobutyric acid

<220>
<221> MOD_RES
<222> (10)..(10)
<223> AMIDATION

<400> 22

Xaa Val Xaa Glu Ile Gln Leu Met His Gln
1 5 10

<210> 23
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> [Ac5c-1, Trp-2, Aib-3, M]PTH(1-14)

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> Xaa represents 1-aminocyclopentane-1-carboxylic acid

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa represents alpha-aminoisobutyric acid

<220>
<221> MISC_FEATURE
<222> (11)..(11)
<223> Xaa represents homoarginine

<220>
<221> MOD_RES
<222> (14)..(14)
<223> AMIDATION

<400> 23

Xaa Trp Xaa Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp
1 5 10

<210> 24
<211> 14
<212> PRT
<213> Artificial Sequence

<220>

-15-

<223> [Deg-1, M] PTH(1-14)

<220>

<221> MISC_FEATURE

<222> (1)..(1)

<223> Xaa represents alpha, alpha-diethylglycine

<220>

<221> MISC_FEATURE

<222> (11)..(11)

<223> Xaa represents homoarginine

<400> 24

Xaa Val Ala Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp
1 5 10

<210> 25

<211> 14

<212> PRT

<213> Artificial Sequence

<220>

<223> [Ac3c-1, M] PTH(1-14)

<220>

<221> MISC_FEATURE

<222> (1)..(1)

<223> Xaa represents 1-aminocyclopropane-1-carboxylic acid

<220>

<221> MISC_FEATURE

<222> (11)..(11)

<223> Xaa represents homoarginine

<220>

<221> MOD_RES

<222> (14)..(14)

<223> AMIDATION

<400> 25

Xaa Val Ala Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp
1 5 10

<210> 26

<211> 14

<212> PRT

<213> Homo sapiens

<220>

<221> MOD_RES

<222> (14)..(14)

<223> AMIDATION

<400> 26

Ser Val Ser Glu Ile Gln Leu Met His Asn Leu Gly Lys His

-16-

1

5

10

<210> 27
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> [Deg-1,3, M]PTH(1-14)

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> Xaa represents alpha, alpha-diethylglycine

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa represents alpha, alpha-diethylglycine

<220>
<221> MISC_FEATURE
<222> (11)..(11)
<223> Xaa represents homoarginine

<220>
<221> MOD_RES
<222> (14)..(14)
<223> AMIDATION

<400> 27

Xaa Val Xaa Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp
1 5 10

<210> 28
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> [Ac3c-1,3, M]PTH(1-14)

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> Xaa represents 1-aminocyclopropane-1-carboxylic acid

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa represents 1-aminocyclopropane-1-carboxylic acid

<220>
<221> MISC_FEATURE
<222> (11)..(11)
<223> Xaa represents homoarginine

-17-

<220>
<221> MOD_RES
<222> (14)..(14)
<223> AMIDATION

<400> 28

Xaa Val Xaa Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp
1 5 10

<210> 29
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> [Ac5c-1,3, M]PTH(1-14)

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> Xaa represents 1-aminocyclopentane-1-carboxylic acid

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa represents 1-aminocyclopentane-1-carboxylic acid

<220>
<221> MISC_FEATURE
<222> (11)..(11)
<223> Xaa represents homoarginine

<220>
<221> MOD_RES
<222> (14)..(14)
<223> AMIDATION

<400> 29

Xaa Val Xaa Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp
1 5 10

<210> 30
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> [Ac5c-1, Aib-3, Gln-10, Ala-12, Trp-14]PTH(1-14)

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> Xaa represents 1-aminocyclopentane-1-carboxylic acid

<220>
<221> MISC_FEATURE

-18-

<222> (3)..(3)
<223> Xaa represents alpha-aminoisobutyric acid

<220>
<221> MOD_RES
<222> (14)..(14)
<223> AMIDATION

<400> 30

Xaa Val Xaa Glu Ile Gln Leu Met His Gln Arg Ala Lys Trp
1 5 10

<210> 31
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> [Ac5c-1, Aib-3, M]PTH(1-9)

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> Xaa represents 1-aminocyclopentane-1-carboxylic acid

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa represents alpha-aminoisobutyric acid

<220>
<221> MOD_RES
<222> (9)..(9)
<223> AMIDATION

<400> 31

Xaa Val Xaa Glu Ile Gln Leu Met His
1 5

<210> 32
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> [Aib-1, Ac5c-3]PTH(1-14)

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> Xaa represents alpha-aminoisobutyric acid

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa represents 1-aminocyclopentane-1-carboxylic acid

-19-

<220>
<221> MOD_RES
<222> (14)..(14)
<223> AMIDATION

<400> 32

Xaa Val Xaa Glu Ile Gln Leu Met His Asn Leu Gly Lys His
1 5 10

<210> 33
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> native hPTH(1-9)amide

<220>
<221> MOD_RES
<222> (9)..(9)
<223> AMIDATION

<400> 33

Ser Val Ser Glu Ile Gln Leu Met His
1 5

<210> 34
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> [Aib-1,3]PTH(1-9)amide

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> Xaa represents alpha-aminoisobutyric acid

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa represents alpha-aminoisobutyric acid

<220>
<221> MOD_RES
<222> (9)..(9)
<223> AMIDATION

<400> 34

Xaa Val Xaa Glu Ile Gln Leu Met His
1 5

<210> 35

-20-

<211> 10
<212> PRT
<213> Artificial Sequence

<220>
<223> [Aib-1,3, Gln-10]PTH(1-10)amide

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> Xaa represents alpha-aminoisobutyric acid

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa represents alpha-aminoisobutyric acid

<220>
<221> MOD_RES
<222> (10)..(10)
<223> AMIDATION

<400> 35

Xaa Val Xaa Glu Ile Gln Leu Met His Gln
1 5 10

<210> 36
<211> 21
<212> PRT
<213> Artificial Sequence

<220>
<223> [Aib-1,3, Nle-8, Gln-10, Har-11, Ala-12, Trp-14, Arg-19,
Tyr-21]rPTH(1-21)amide radiolabeled with Iodine isotope 125

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> Xaa represents alpha-aminoisobutyric acid

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa represents alpha-aminoisobutyric acid

<220>
<221> MISC_FEATURE
<222> (8)..(8)
<223> Xaa represents norleucine

<220>
<221> MISC_FEATURE
<222> (11)..(11)
<223> Xaa represents homoarginine

<220>
<221> MOD_RES
<222> (21)..(21)

-21-

<223> AMIDATION

<220>

<221> MISC_FEATURE

<222> (21)..(21)

<223> Radiolabeled with Iodine isotope 125

<400> 36

Xaa Val Xaa Glu Ile Gln Leu Xaa His Gln Xaa Ala Lys Trp Leu Ala
1 5 10 15

Ser Val Arg Arg Tyr
20

<210> 37

<211> 14

<212> PRT

<213> Artificial Sequence

<220>

<223> [Deg-3, M]PTH(1-14)

<220>

<221> MISC_FEATURE

<222> (3)..(3)

<223> Xaa represents alpha, alpha-diethylglycine

<220>

<221> MISC_FEATURE

<222> (11)..(11)

<223> Xaa represents homoarginine

<220>

<221> MOD_RES

<222> (14)..(14)

<223> AMIDATION

<400> 37

Ala Val Xaa Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp
1 5 10

<210> 38

<211> 14

<212> PRT

<213> Artificial Sequence

<220>

<223> [Ac3c-3, M]PTH(1-14)

<220>

<221> MISC_FEATURE

<222> (3)..(3)

<223> Xaa represents 1-aminocyclopropane-1-carboxylic acid

<220>

-22-

<221> MISC_FEATURE
<222> (11)..(11)
<223> Xaa represents homoarginine

<220>
<221> MOD_RES
<222> (14)..(14)
<223> AMIDATION

<400> 38

Ala Val Xaa Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp
1 5 10

<210> 39
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> [Ac5c-3, M]PTH(1-14)

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa represents 1-aminocyclopentane-1-carboxylic acid

<220>
<221> MISC_FEATURE
<222> (11)..(11)
<223> Xaa represents homoarginine

<220>
<221> MOD_RES
<222> (14)..(14)
<223> AMIDATION

<400> 39

Ala Val Xaa Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp
1 5 10

<210> 40
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> [Deg-1, Ac3c-3, M]PTH(1-14)

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> Xaa represents alpha, alpha-diethylglycine

<220>
<221> MISC_FEATURE
<222> (3)..(3)

-23-

<223> Xaa represents 1-aminocyclopropane-1-carboxylic acid

<220>

<221> MISC_FEATURE

<222> (11)..(11)

<223> Xaa represents homoarginine

<220>

<221> MOD_RES

<222> (14)..(14)

<223> AMIDATION

<400> 40

Xaa Val Xaa Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp
1 5 10

<210> 41

<211> 14

<212> PRT

<213> Artificial Sequence

<220>

<223> [Deg-1, Ac5c-3, M]PTH(1-14)

<220>

<221> MISC_FEATURE

<222> (1)..(1)

<223> Xaa represents alpha, alpha-diethylglycine

<220>

<221> MISC_FEATURE

<222> (3)..(3)

<223> Xaa represents 1-aminocyclopentane-1-carboxylic acid

<220>

<221> MISC_FEATURE

<222> (11)..(11)

<223> Xaa represents homoarginine

<220>

<221> MOD_RES

<222> (14)..(14)

<223> AMIDATION

<400> 41

Xaa Val Xaa Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp
1 5 10

<210> 42

<211> 14

<212> PRT

<213> Artificial Sequence

<220>

<223> [Deg-1, Alb-3, M]PTH(1-14)

-24-

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> Xaa represents alpha, alpha-diethylglycine

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa represents alpha-aminoisobutyric acid

<220>
<221> MISC_FEATURE
<222> (11)..(11)
<223> Xaa represents homoarginine

<220>
<221> MOD_RES
<222> (14)..(14)
<223> AMIDATION

<400> 42

Xaa Val Xaa Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp
1 5 10

<210> 43
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> [Ac3c-1, Deg-3, M]PTH(1-14)

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> Xaa represents 1-aminocyclopropane-1-carboxylic acid

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa represents alpha, alpha-diethylglycine

<220>
<221> MISC_FEATURE
<222> (11)..(11)
<223> Xaa represents homoarginine

<220>
<221> MOD_RES
<222> (14)..(14)
<223> AMIDATION

<400> 43

Xaa Val Xaa Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp
1 5 10

<210> 44

-25-

<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> [Ac3c-1, Ac5c-3, M] PTH(1-14)

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> Xaa represents 1-amino-cyclopropane-1-carboxylic acid

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa represents 1-amino-cyclopentane-1-carboxylic acid

<220>
<221> MISC_FEATURE
<222> (11)..(11)
<223> Xaa represents homoarginine

<220>
<221> MOD_RES
<222> (14)..(14)
<223> AMIDATION

<400> 44

Xaa Val Xaa Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp
1 5 10

<210> 45
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> [Ac3c-1, Aib-3, M] PTH(1-14)

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> Xaa represents 1-aminocyclopropane-1-carboxylic acid

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa represents alpha-aminoisobutyric acid

<220>
<221> MISC_FEATURE
<222> (11)..(11)
<223> Xaa represents homoarginine

<220>
<221> MOD_RES
<222> (14)..(14)
<223> AMIDATION

-26-

<400> 45

Xaa Val Xaa Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp
1 5 10

<210> 46

<211> 14

<212> PRT

<213> Artificial Sequence

<220>

<223> [Ac5c-1, Deg-3, M]PTH(1-14)

<220>

<221> MISC_FEATURE

<222> (1)..(1)

<223> Xaa represents 1-aminocyclopentane-1-carboxylic acid

<220>

<221> MISC_FEATURE

<222> (3)..(3)

<223> Xaa represents alpha, alpha-diethylglycine

<220>

<221> MISC_FEATURE

<222> (11)..(11)

<223> Xaa represents homoarginine

<220>

<221> MISC_FEATURE

<222> (51)..(51)

<223> Xaa represents homoarginine

<400> 46

Xaa Val Xaa Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp
1 5 10

<210> 47

<211> 14

<212> PRT

<213> Artificial Sequence

<220>

<223> [Ac5c-1, Ac3c-3, M]PTH(1-14)

<220>

<221> MISC_FEATURE

<222> (1)..(1)

<223> Xaa represents 1-aminocyclopentane-1-carboxylic acid

<220>

<221> MISC_FEATURE

<222> (3)..(3)

<223> Xaa represents 1-aminocyclopropane-1-carboxylic acid

<220>

<221> MISC_FEATURE

-27-

<222> (11)..(11)
<223> Xaa represents homoarginine

<220>
<221> MOD_RES
<222> (14)..(14)
<223> AMIDATION

<400> 47

Xaa Val Xaa Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp
1 5 10

<210> 48
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> [Aib-1, Deg-3, M]PTH(1-14)

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> Xaa represents alpha-aminoisobutyric acid

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa represents alpha, alpha-diethylglycine

<220>
<221> MISC_FEATURE
<222> (11)..(11)
<223> Xaa represents homoarginine

<220>
<221> MOD_RES
<222> (14)..(14)
<223> AMIDATION

<400> 48

Xaa Val Xaa Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp
1 5 10

<210> 49
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> [Aib-1, Ac3c-3, M]PTH(1-14)

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> Xaa represents alpha-aminoisobutyric acid

-28-

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa represents 1-aminocyclopropane-1-carboxylic acid

<220>
<221> MISC_FEATURE
<222> (11)..(11)
<223> Xaa represents homoarginine

<220>
<221> MOD_RES
<222> (14)..(14)
<223> AMIDATION

<400> 49

Xaa Val Xaa Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp
1 5 10

<210> 50
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<223> [Aib-1, Ac5c-3]PTH(1-14)

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> Xaa represents alpha-aminosiobutyric acid

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa represents 1-aminocyclopentane-1-carboxylic acid

<220>
<221> MISC_FEATURE
<222> (11)..(11)
<223> Xaa represents homoarginine

<220>
<221> MOD_RES
<222> (14)..(14)
<223> AMIDATION

<400> 50

Xaa Val Xaa Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp
1 5 10

<210> 51
<211> 14
<212> PRT
<213> Artificial Sequence

<220>

-29-

<223> [Ac5c-1, Gln-10, Har-11, Ala-12, Trp-14]PTH(1-14)

<220>

<221> MISC_FEATURE

<222> (1)..(1)

<223> Xaa represents 1-aminocyclopentane-1-carboxylic acid

<220>

<221> MISC_FEATURE

<222> (11)..(11)

<223> Xaa represents homoarginine

<220>

<221> MOD_RES

<222> (14)..(14)

<223> AMIDATION

<400> 51

Xaa Val Ser Glu Ile Gln Leu Met His Gln Xaa Ala Lys Trp
1 5 10

<210> 52

<211> 14

<212> PRT

<213> Artificial Sequence

<220>

<223> [Ac5c-1]PTH(1-14)

<220>

<221> MISC_FEATURE

<222> (1)..(1)

<223> Xaa represents 1-aminocyclopentane-1-carboxylic acid

<220>

<221> MOD_RES

<222> (14)..(14)

<223> AMIDATION

<400> 52

Xaa Val Ser Glu Ile Gln Leu Met His Asn Leu Gly Lys His
1 5 10

2. STATEMENT CONCERNING NON-PREJUDICIAL DISCLOSURES OR EXCEPTIONS TO LACK OF NOVELTY

Due to a disclosure by the inventors on 22 September 2002, the applicant respectfully requests that the subject International application be granted the respective provisions under National laws concerning Exceptions to Lack of Novelty in each of the designated countries. This is not an admission that the subject invention lacks novelty or inventive step over this disclosure. Exception to Lack of Novelty is hereby requested for purposes of disclosure and precautionary measures.

INTERNATIONAL SEARCH REPORT

International application No.

PCT/US03/08261

A. CLASSIFICATION OF SUBJECT MATTER

IPC(7) : A61K 38/29; C07K 1/06, 14/635

US CL : 514/14

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

U.S. : 514/14 , 2; 530/387.2, 387.9, 388.24, 387.1

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)

Please See Continuation Sheet

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	WO 97/02839 A1 (BIOMEASURE INCORPORATED) 30 January 1997 (30.01.1997), see entire document.	1, 26
---		-----
Y		2-25, 27-32
A	US 6,030,790 A (ADERMANN et al) 29 February 2000 (29.02.2000), see entire document.	1-49
A	US 2002/0110871 A1 (ZAHRADNIK et al) 15 August 2002 (15.08.2002), see entire document.	1-49
A	SHIMIZU et al. Enhanced Activity in Parathyroid Hormone-(1-14) and -(1-11): Novel Peptides for Probing Ligand-Receptor Interactions. Endocrinology 2001, Vol. 142, No. 7, pages 3068-3074, entire document.	1-49
A	CHOREV et al. Modifications of Position 12 in Parathyroid Hormone and Parathyroid Hormone Related Protein: Toward the Design of Highly Potent Antagonists. Biochemistry 1990, Vol. 29, pages 1580-1586, see entire document.	1-49

☐ Further documents are listed in the continuation of Box C.

☐ See patent family annex.

* Special categories of cited documents:

"A" document defining the general state of the art which is not considered to be of particular relevance

"E" earlier application or patent published on or after the international filing date

"L" document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)

"O" document referring to an oral disclosure, use, exhibition or other means

"P" document published prior to the international filing date but later than the priority date claimed

"T" later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention

"X" document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone

"Y" document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art

"&" document member of the same patent family

Date of the actual completion of the international search

10 June 2003 (10.06.2003)

Date of mailing of the international search report

24 OCT 2003

Name and mailing address of the ISA/US

Mail Stop PCT, Attn: ISA/US
Commissioner for Patents
P.O. Box 1450
Alexandria, Virginia 22313-1450

Facsimile No. (703)305-3230

Authorized officer

B. Dell Chism

Telephone No. (703)308-0196

INTERNATIONAL SEARCH REPORT

PCT/US03/08261

Continuation of B. FIELDS SEARCHED Item 3:

STN REGISTRY CAPLUS USPATFULL WEST BIOSIS MEDLINE SCISEARCH

search terms: parathyroid, hormone, PTH, analogs, Aib, diethylglycine, cyclohexylalanine, Cha

**This Page is Inserted by IFW Indexing and Scanning
Operations and is not part of the Official Record**

BEST AVAILABLE IMAGES

Defective images within this document are accurate representations of the original documents submitted by the applicant.

Defects in the images include but are not limited to the items checked:

- ☐ **BLACK BORDERS**
- ☐ **IMAGE CUT OFF AT TOP, BOTTOM OR SIDES**
- ☐ **FADED TEXT OR DRAWING**
- ☐ **BLURRED OR ILLEGIBLE TEXT OR DRAWING**
- ☐ **SKEWED/SLANTED IMAGES**
- ☐ **COLOR OR BLACK AND WHITE PHOTOGRAPHS**
- ☐ **GRAY SCALE DOCUMENTS**
- ☐ **LINES OR MARKS ON ORIGINAL DOCUMENT**
- ☐ **REFERENCE(S) OR EXHIBIT(S) SUBMITTED ARE POOR QUALITY**
- ☐ **OTHER:** _____

IMAGES ARE BEST AVAILABLE COPY.

As rescanning these documents will not correct the image problems checked, please do not report these problems to the IFW Image Problem Mailbox.